

SEQUENCE LISTING

<110> LEVINE, et al.

<120> VARIANTS OF PROTEIN KINASES

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<150> 09/724,676

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<150> 136776

<151> 2000-06-15

<150> 135619

<151> 2000-04-12

<160> 182

<170> PatentIn version 3.0

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<212> DNA
<213> Homo sapiens

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aaggcaactt gttcacccctg gtgcctccca gccgctccct gagcacaat ggcgagaaca	1140
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agtgtctgaa gcctggctcc actattgtgt gacaagcctc ttctcctctc tgaatcttta	1440
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tttagggggc tttggaaggg aaatgcatt gaatcaatgc angaagaagc attaaccaa	1560
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ggcactgggg cttgctccag attgtggtgg gagaatgctc tactaagaga ttgatgggt	1680
gctgggggtgg agggggggaa gcctgnagcc caagagaccc tgttcctggn agaatgaatg	1740
ggaaatattc ataaataatg tacacaaagt aactcttcc ttctgctctc cctgttagct	1800

cccaagtgcc ccccatcaaa cctgggcgcc cg 1832

<210> 21
<211> 1269
<212> DNA
<213> Homo sapiens

<400> 21
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gtggagtgcc cttttgtga tgaagttcc aaatacgaga agctcgccaa gatcgccaa 180
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aagaagggtgc tgatggaaaa cgagaaggag gggttccccca ttacagcctt gcgggagatc 300
aagatccctt agcttctaaa acacgagaat gtggtaact tgattgagat ttgtcgaacc 360
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ttggcaccac cgccggaa gggcagccag atcaccacg agtccaccaa ccagagtcgc 660
aatccgcac ccaccaacca gacggagttt gagcgcgtct tctgaggccc ggcgcttgcc 720
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tgtccttgct ggtttctgg atggttccccca gagggttcc atggggtagg aggatggct 960
cgccaccag tgacttttc taagagctcc cggcgctggc gaagagggga caggtccctc 1020
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gaatgggctg ttttggctt aaccctcaga aacactgggg ctggcacaaa ctcttggttt 1140
cttcaacagg agaattttac tgtgtttttt ttgggtccat tgtttggaga cattcctggg 1200
cacagtttgg tccgttagaa ttaaaaagttt aattttttttt tttttttttt 1260
yccccccaaa 1269

<210> 22
<211> 623

<212> DNA
<213> Homo sapiens

<220>
<221> -
<222> (1)..(623)
<223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

<400> 22
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gggggcctct ctagcttgcg gcctgtgtct atggtcgggc cctctgcgtc cagctgtctcc 120
ggaccgagct cgggtgtatg gggccgtagg aaccggctcc ggggccccga taacgggccc 180
cccccacago accccgggct ggcgtgaggg tctcccttga tctgagaatg gctacctctc 240
gatatgagcc agtggctgaa attggtgtcg gtgcctatgg gacagtgtac aaggcccgtg 300
atccccacag tggccacttt gtgcctcaa gagtgtgaga gtccccaatg gaggaggagg 360
tggaggaggo ctcccatca gcacagttcg tgaggtggct ttactgaggc gactggaggc 420
tttgagcat cccaatgttg tccggctgat ggacgtctgt gccacatccc gaactgaccg 480
ggagatcaag gtaaccctgg tggggatca tgtagaccag gacctaagga catactggaa 540
caaggcaccc ccaccaggct tgccagccga aacgatcaag gtgagtgaaa ttggtaggca 600
ttganaggtg gattgggacc ttt 623

<210> 23
<211> 502
<212> DNA
<213> Homo sapiens

<400> 23
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ggaccgagct cgggtgtatg gggccgtagg aaccggctcc ggggccccga taacgggccc 180
cccccacago accccgggct ggcgtgaggg tctcccttga tctgagaatg gctacctctc 240
gatatgagcc agtggctgaa attggtgtcg gtgcctatgg gacagtgtac aaggcccgtg 300
atccccacag tggccacttt gtgcctcaa gagtgtgaga gtccccaatc acctctcatt 360
ttgaggcttc tccttcctt tcccatattt ctacactaag ggttatgttc cctttgtcc 420
cttccctac ctttatattt ggggtccttt tttatacagg aaaaacaaaa caaagaaata 480
aagtgcacgc ggccgcgaat tc 502

<210> 24
 <211> 1148
 <212> DNA
 <213> Homo sapiens

<220>
 <221> -
 <222> (1)..(1148)
 <223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

<400> 24	
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tcacttgcnt ttgagccaca caccctccag ccactaccca tagcagttat ggggtgtgt	120
cctgggaaga ctcctttctg ctccccaaaa ccccaaggcc tggctcgggg ccactggagc	180
cgcaggcggg acatatgtgt gaccggccct ctgcccctgg cagccccgcg ctgtgtactg	240
taaggacgtg ctggacatcg agcagttctc cactgtgaag ggcgtcaatc tggaccacac	300
agacgacgac ttctactcca agttctccac gggctctgtg tccatcccat ggcaaaacga	360
gatgatagaa acagaatgct ttaaggagct gaacgtgttt ggacctaatg gtaccctccc	420
gccagatctg aacagaaaacc accctccgga accgccccaaag aaagggctgc tccagagact	480
cttcaagcgg cagcatcaga acaattccaa gagttcgccc agctccaaga ccagtttaa	540
ccaccacata aactcaaacc atgtcagctc gaactccacc ggaagcagct agttcggct	600
ctggcctcca agtccacagt ggaaccagcc cagacccttc tccttagaag tggaagtagt	660
ggagccccctg ctctggtggg gctgccaggg gagaccccg gагccggggaggaggccgt	720
ccatccccgtc gacgtagaac ctcgaggaaa ctcaaaagaaa tttccactca ggtctgttt	780
ccgaggcggc cccggccggg gtggattgga tttgtctttg gtgaacatttca caatagaaat	840
ccaattggat acgacaactt gcacgtattt taatagcgtc ataactagaa ctgaattttt	900
tctttatgtat tttaaaagaa aagttttgtaa aatttctcta ctgtctcagt ttacattttt	960
tatatttgtat tttaaaatgaa agtgagactt tgagggtgtat tttttctgt gcagccactg	1020
ttaagccatg tgttccaagg cattttagcg gggaggggggt tatcaaaaaaa aaaaaaaatgt	1080
gactcaagac ttccagagcc tcaaataatgaga aaatgtcttt attaaatgtaa gaaagtgtatc	1140
cataacttc	1148

<210> 25
 <211> 1679
 <212> DNA

<213> Homo sapiens

<220>

<221> -

<222> (1)..(1679)

<223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

<400> 25

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ggcattttcc tgtgtccctcc cacacctgca ctctccaaag gctttgtgg tcactctgag 180

atggcagatt ggggtccctg ttgtccctgg acagatgaga atgccgagag ctcgtatgcc 240

tngccaagg gcacacagca aggccgggtg cccatgcggc tgcaggagg acccaactgac 300

cctgctgtcc ccctcaggcc acattaggat ctcagacctg ggcttggctg tgaagatccc 360

cgagggagac ctgatccgcg gccgggtggg cactgttggc tacatggctc cagaggtcct 420

gaacaaccag aggtacggcc tgagccccga ctactggggc cttggctgcc tcatactatga 480

gatgatcgag ggccagtcgc cgttccgcgg ccgcaaggag aaggtgaagc gggaggaggt 540

ggaccgcgg gtcctggaga cggaggaggt gtactcccac aagttctccg aggaggccaa 600

gtccatctgc aagatgctgc tcacgaaaga tgcgaagcag aggctggct gccaggagga 660

gggggctgca gaggtcaaga gacacccctt cttaggaac atgaacttca agcgctt 720

agccgggatg ttggaccctc cttcggttcc agaccccccgc gctgtgtact gtaaggacgt 780

gctggacatc gagcaggcttccactgtgaa gggcgtaat ctggaccaca cagacgacga 840

cttctactcc aagttotcca cgggctctgt gtccatccca tggcaaaacg agatgataga 900

aacagaatgc tttaaggagc tgaacgtgtt tggacctaattt ggtaccctcc cgccagatct 960

gaacagaaac caccctccgg aaccgcggaa gaaagggtg ctccagagac tcttcaagcg 1020

gcagcatcag aacaattcca agagttcgcc cagctccaaag accagttta accaccacat 1080

aaactcaaac catgtcagct cgaactccac cgaaagcagc tagttcggc tctggcctcc 1140

aagtccacag tggaaaccagc ccagaccctt ctccttagaa gtggaaagttag tggagcccct 1200

gctctgggtgg ggctggccagg ggagaccccg ggagccgggg aaggaggccg tccatcccgt 1260

cgacgtagaa cctcgagggtt tctcaaagaa atttccactc aggtctgttt tccgaggcgg 1320

ccccggccgg ggtggattgg atttgtcttt ggtgaacatt gcaatagaaa tccaatttgg 1380

tacgacaact tgcacgtatt ttaatagcgt cataactaga actgaatttt gtctttatga 1440

tttttaaaga aaagtttgt aaatttctct actgtctcag tttacatttt gtatatttgt 1500
attnaaatga aagtgagact ttgagggtgt atatttctg tgcagccact gttaagccat 1560
gtgttccaag gcattttagc ggggaggggg ttatcaaaaa aaaaaaaaaatg tgactcaaga 1620
cttccagagc ctcaaatgag aaaatgtctt tattaaatgt agaaaagtgat ccatacttc 1679

<210> 26
<211> 897
<212> DNA
<213> Homo sapiens

<220>
<221> -
<222> (1)..(897)
<223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

<400> 26
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ggcattttcc tgtgtccctcc cacacctgca ctctcccaag gctttgtgg tcactctgag 180
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cctgctgtcc ccctcaggcc acattaggat ctcagacctg ggcttggctg tgaagatccc 360
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gaacaaccag aggtacggcc tgagccccga ctactggggc cttggctgcc tcatactatga 480
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cctccctggg cctggcccca gtcgtcccc agaacagcaa acaggctgaa gggacagggg 720
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aggactgggg tggtcagcgg agacttccag gagaaagcct ggggtggggca gggacaccca 840
gtaaccaagg gaagagggggt gaggggaaca gtgtggaaca ccctcacatc cagcctg 897

<210> 27
<211> 1224
<212> DNA
<213> Homo sapiens

<400> 27
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cgaggcgaac cagcccttct gtgccgtgaa gatgaaggag gcgcctcagca cagagcgtgg 180
gaaaacactg gtgcagaaga agccgaccat gtatcctgag tggaagtoga cgttcgatgc 240
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gtctgaggtg accgtgggtg tgtcggtgct ggccgagcgc tgcaagaaga acaatggcaa 360
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cctggaggac gtggattgca aacagtctat gcgcagttag gacgaggcca agttcccaac 480
gatgaaccgc cgccggagcca tcaaacadggc caaaatccac tacatcaaga accatgagtt 540
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cctcaacaag caaggotaca aatgcaggca atgtaacgct gccatccaca agaaatgcat 660
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gaaagaacgc ttcaacatcg acatgccgca ccgcctcaag gttcacaact acatgagccc 780
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gagaaaaacc atggggact ggtgatatgt tgatctttt caaaaaata tatatatgac 1140
aaaaaaaaaa aaaaaaggag cacaagctgt ttgaaccacc agtttattt gtgtgtctaa 1200
ataaacacca aatagtacca aaaa 1224

<210> 28
<211> 1424
<212> DNA
<213> Homo sapiens

<400> 28
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aggttagggtt ggaggaggca aatgatatgg tggaacaagg accttgaaat caatccaaaa 180
cccaggtttt ccttaggaagg ccacccggaa cccatggtaa gccaactgtt gcgcaggat 240

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cctggcagcg	ttgaggaggc	tccagcagag	atccagtgt	aagggcccta	gaggggctgg	360
tggcatcccc	tcaatcttgg	tcctctctcc	ccagagttgt	gtgggacccc	aggtatcta	420
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aagggttttgc	atcattccag	ctcctctggg	ctctggcctc	aggcccacta	atgatcctgc	1260
taccctcttg	aagaccagcc	cggtacctct	ctccccactg	gccaggactc	tgagatcaga	1320
gctgggttgg	aaggagcca	ttctgaacgc	cacgcctggc	ccggtcagtg	ctgcatgcac	1380
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<210> 29
 <211> 2027
 <212> DNA
 <213> Homo sapiens

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gcctagcgag	cctgcgggtc	gaccccagcc	agcgcagcga	cggggcgctg	cctggcccag	180
gcgcacacgg	aagtgcgttt	ctctgaagta	gctttggaaa	gtagagaaga	aaatccagtt	240
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gttcgtctcc ctgtgcccaga gccaaggctgt atccccatcaq taatgtgctq agaccccaat 1980

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2027

<210> 30
<211> 1609
<212> DNA
<213> Homo sapiens

<400> 30
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gcccggccccc gcgccgcacc gcccgggccc gccgccccgg ccagggaggg attcggccgc 300
cggggccgggg acaccccgcc gccgccccct cgggtgcgttc ggaaggccca ccggctcccg 360
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catgtgagtgc ggcgcggag cctcagcgcc gcgcagttttt ttgttgaagaag caggatgtgc 480
atctaaacgt ggaaaaagac cagtcctgccc tctgttgtag aagacatgtg gtgttatataa 540
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ataaagaagc aacaaaactg acggaggaga gggacggcag cctgaaccag agctctgggt 660
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<210> 31
<211> 1995
<212> DNA
<213> Homo sapiens

<220>
<221> -
<222> (1)..(1995)
<223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

<400> 31
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cgggccgggg acaccccgcc gccgcgcct cggtgcttc ggaaggccca ccggctcccg 360
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catgtgagtg ggctccggag cctcagcgcc gcgcagtttt ttgttgcgatgt 480
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atcatggctc actacaatct ctgccttcct gcctcaaggg ttcttccttg agcacccatcg 660
cctcccaagt agctggacc acaggaattt agataatggg ctgtgtgcaa tgtaaggata 720
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aaccaccagt tatgtgctcg ttaatgaaaa catttttaa aacagactaa cttgcgttt 1920
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<210> 32
<211> 2590
<212> DNA
<213> Homo sapiens

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cccgcttacg tatggaaagt gtctttgtct ctccctggagc tgcacaagag gagaaagcc 180
ctgactgagc ctgaggccccg atactaccta cggcaaattg tgcttggctg ccagtacctg 240
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cacagttcg aggtggatgt gtggccatt gggtgtatca tgtatacctt gtttagtggc 480
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actttctcag 2590

<210> 33
<211> 1096
<212> DNA
<213> Homo sapiens

<220>
<221> -
<222> (1)..(1096)
<223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

<400> 33
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ggaaggccat ggctctgttg ctccttttc ttgcctctca cagattggaa gtatctaggg 180
acagtgggtg gctaggacag tgctggctgc aggggtctg ggagcgtkgg ctcacagtg 240
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acgggtccca ggacagactc atagctagac cccgttggcg gcctctgtgt tgaaccagaa 900
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ccaccctgct agatggtag gaccaaggct ctgttctcct ggaagccaaag gtccggcacat 1020

DRAFT

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ggcagagaat gggggg	1096

<210> 34
<211> 940
<212> DNA
<213> Homo sapiens

<400> 34	
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acagtgggtg gctaggacag tgctggctgc aggggtctg ggagcgkkg cctcacagtg	240
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gctccctgcg gtttgcac aagcccact acttttacct acgtcagctc ttccgcaacc	360
tcttccaccg gcagggcttc tcctatgact acgtcttga ctggaacatg ctgaaattcg	420
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gtccccgccc ctgtccctgt gctgggccc cgtactcacc cacgtactgg taaggatcct	540
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aaaaggccct gccatgggt ctttggggc caggacagcc tgagggaggg atggtggcca	660
ctgcccacaa gggccttgtt gggAACGGGT CCCAGGACAG ACTCATAGCT AGACCCCGTT	720
ggcggcctct gtgttgaacc agaactcatt aaacacctcc tcttgcttca aaawrgtgtg	780
cctctttcat ggcaggcccc tcagccaccc tgcttagatgg ttaggaccaa ggctctgttc	840
tccttggaaac caaggtcgcc acattggtct tgggcttctc ttctctctgg gtttcttgg	900
cactaaggag taacacagag gtcaggcaga gaatgggggg	940

<210> 35
<211> 951
<212> DNA
<213> Homo sapiens

<220>
<221> -
<222> (1)..(951)
<223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

<400> 35

ggctcctgga tccggctacc tggcagaggc tcctggccac tgttcgagg tcttcccttg 60
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ggaaggccat ggctctgttg ctccttttc ttgcctctca cagattgaa gtatctaggg 180
acagtgggtg gctaggacag tgctggctgc agggggctg ggagcgtkgg ctcacagtg 240
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gctccctgctg 6ttgacgac aagcccact actttacct acgtcagctc ttccgcaacc 360
tcttccaccg gcagggcttc tcctatgact acgtcttga ctgaaacatg ctgaaattcg 420
gcggccccct tcctgcagc cccctgcct tccctgtgga cggccccagg atgaactagg 480
gtcagccccg gaatcccgag gatgtggacc gggagcggcg agaacacgaa cgcgaggaga 540
ggatggggca gctacggggg tccgcgaccc gagccctgccc ccctggccca cccacggggg 600
ccactgccaa cccgcctccgc agtgcgcggc agccctgtgc ttccacgcca gcctcccgca 660
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ggcaagaggt ctcccgatc ccagcctcac agacaagtgt gccatttgac catctcgga 840
agtgaggaga gccccattg gaccagtgtt tgcttagtgtt cttcaactgtttaa 900
aaamaaaaam aaaaawaaam srcmaaaakw acmacwmaaa aacccagcac a 951

<210> 36
<211> 2063
<212> DNA
<213> Homo sapiens

<220>
<221> -
<222> (1)..(2063)
<223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

<400> 36
gcggccgccc gaaaagcgct tcggccacat ccagcagcag tagcagccgc aaggaccggg 60
actcgaaggc ccaccgcagc cggactaagt cgtccaagga gccccttcg gcctacaagg 120
aaccgccccaa ggcctaccgg gaggacaaga ccgagcctaa ggcctacagg cggccggcggt 180
ccctcagccc actgggaggc cgggacgaca gcccgggtgc ccacaggccc tctcagagcc 240
tgaggagccg caagtcccc agcccgccag gaggtggcag cagcccstat tctcggccgc 300
tgccgcgctc cccgagcccc tacagtcgcc gcccgtcccc cagctacagc cgccacagct 360

cctacgagcg gggcggcgac gtgtccccta gtcctacag cagcagcgc tggcgccgct 420
ctcgcagtcc ctacagccct gtgctcagac ggtctggaaa atcccgaagc agaagcccgt 480
attcatctag gcattcaaga tctcgttagca ggcacagatt gtctagatcc agaagtcgtc 540
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agaataaaaa agcacgagca gcagaggcag caagagccgc agaagcagcg aaagctgcag 660
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<210> 37
<211> 1393
<212> DNA
<213> Homo sapiens

<400> 37
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gagggcgcgg ccgcgggcgg ggctgagggc ggccggggcgg ggccgcccga gctgggaggg 120
ccgcggcgcc gaggggagga gagcggccca tggacccgcg gggcccgccg ccccagactc 180
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ggggctgttt ttc 1393

<210> 38
<211> 1244
<212> DNA
<213> Homo sapiens

<400> 38
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gagggcgcgg ccgcggcgg ggctgagggc ggccccggcg ggccgcccga gctgggaggg 120
ccgcggcgcc gaggggagga gagcggccca tggaccccgcg gggcccgcg ccccaagactc 180
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cgcaacgaga gcgcctgag cgtggaaacg ctgcgcacg tgctcgtctg cctgtacacc 480
gagtgcagcc actcggccct gcgcgcgac aagtacgtgg ccgagttcct cgagtggct 540
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<210> 39
<211> 3171
<212> DNA
<213> Homo sapiens

<400> 39
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cagttagact tcaaggaaga gaataagctc gccacataaa gaggaaacaa agagcaaccc 120
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DNA sequence analysis

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<223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

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<212> DNA
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tat	tttatct	ac	ttt tttca gaaagtccctt agtgc	1920
tctagtcaga	a	atgtctgtc agatagttag aattgtaaca tctaaacctg ccacggatcg	1980	
aatggtactt	ac	aggtac	ct ctttaggga ctctgtatc cctaaaatat cagaagaaaa	2040
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<210> 58
 <211> 1097
 <212> DNA
 <213> Homo sapiens

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		gtggggcggc	gcccggctggg cggccgggccc ggagttgcct cccggggccc gcgcgtgaggg	180	
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		gcgtgagtct	ctacctgagc gaggacgagg tgccggcgt gatcggtctt gatgcagaac	360	
		tttattatgt	gagaaatgac cttattatgc actacgctct atcctttagt ctgttagtac	420	
		ccagtgagac	aaatttcctg cacttcaccc ggcacatgc gtc	ccaaagggtt gaatataagc	480
		tgggattcca	agtggacaat gtttggcaa tggatatgcc ccaggtcaac atttctgttc	540	
		agggggaa	gt tccacgcact ttatcagtgt ttccggtaga gctttctgt actggcaaag	600	
		tagattctga	atgttata gata ctaatgcagc tcaacttgac agtaaattct tcaaaaaatt	660	
		ttaccgtctt	aaattttaaa cgaaggaaaaa tgtgctacaa aaaacttgaa gaagtaaaaa	720	
		cttcagcc	ttt ggacaaaaac actagcagaa ctattatga tcctgtacat gcagctccaa	780	
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		tagcaataat	at tagctgtt ttgcacccatc atagatgaa aaggattgaa ctggatgaca	900	
		ggtattgtac	atatttggg aaagaaaaaa aatgaaagca gttatttta tataatgtggg	960	
		agcccataca	catctgatga cagtggtgca gggaaaggag ggaggatgat taagccccag	1020	

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ttttaaaaat taaggtc 1097

<210> 59
<211> 3382
<212> DNA
<213> Homo sapiens

<220>
<221> -
<222> (1)..(3382)
<223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

<400> 59
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gtggggcggcgc gcccgtgggg cggccggggcc ggagttgcct cccggggccc gcgctgaggg 180
ccccggcgcg cccggccgctg ctgtttctgc ttgcgtgtt gcccgtgtg cccggccctg 240
gcccgtgccgc cggccggcc cccggccccc cggagctgca gtcggcttcc gggggccca 300
gcccgtgccgc cggccggcc cccggccccc cggagctgca gtcggcttcc gggggccca 360
tttattatgt gagaatgac cttattatgc actacgctct atccttagt ctgttagtac 420
ccagttagac aaatttcctg cacttcacct ggcacatcgaa gtccaaagggtt gaatataagc 480
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aattaatgtt ataaggaagt attaaaacac tgaaacattt ctccagaacc agcaagtaag 3120
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aatgtgtgc agtngaggc ttttgctgtt tttaaaaaag ccttatgaat cagcagcaca 3300
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ctgtggtaca ttttatttaa ca 3382

<210> 60
<211> 2195
<212> DNA
<213> Homo sapiens

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aagccatgga aggaaaggtg ccaagtggtc aggcgagagc ctccagggca aaggccttgg 180
gcaggtggga atcctgattt gttcctgaaa ggtagttgg ctgaatcatt cctgagaagg 240
ctggagaggc cagcaggaaa caaaacccag caaggccttt tgctgtgagg gcattaggga 300
gctggagggta ttttgagcag cagagggaca taggttgtgt tagtgtttga gcaccagccc 360
tctggcccc ttgttagatt tagaggacca gactcaggga tggggcttga gggaggttag 420
graaggaagg gggcttgga tcattgcagg agctatgggg attccagaaa tggtgaggga 480
acggaggagt agggggtaaa caaggattcc tagcctggaa ccagtgcacc agtcctgagt 540
cttccaggac cacaggcagc cttaagcctg gtccccatac acaggctgaa gtggcagtcc 600
cageggctgt ccctgcggca gaggctgagg ccgaggtgac gctgcggag ctccaggaag 660
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ctagaactgt	cttcgactcc	ggggcccccgt	tggaagactg	agtgcgggg	gcacggcaca	660
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tgtggcccag	ctgtgccacc	gagcgtcgag	aagagggggc	tgggctggca	gcgcgcgcgg	1620
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<210> 62
 <211> 1149
 <212> DNA
 <213> Homo sapiens

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agctgtctta ggcataatat ttgtctgcct ctaattattt ccttcttaggt ttgtacaagt 180
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tgattgcacc aggtttcag gttaagtcta taaataatga agagaggaag gaacatagaa 300
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aaatccttgg agaaagctta ggctgttaac ccagtcactc caccttgac acattactag 600
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cttaaaaata 1149

<210> 63
<211> 1461
<212> DNA
<213> Homo sapiens

<220>
<221> -
<222> (1)..(1461)
<223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

<400> 63
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tccagtcag cgtgggttta tggctgtcc catccccatc aatggtgagg acccagactg 180
agtccagcac gccccctggc attcctggtg gcagcaggca gggccccggcc atggacggca 240

ctgcagccga	gcctcgcccc	ggcgccggct	ccctgcagca	tgcccagcct	ccgcgcgcagc	300
ctcgaagaa	gccccctgag	gacttcaagt	ttggaaaaat	ccttggggaa	ggctttttt	360
ccacgggtgt	cctggctcga	gaactggcaa	cctccagaga	atatgcgatt	aaaattctgg	420
agaagcgaca	tatcataaaa	gagaacaagg	tcccctatgt	aaccagagag	cggatgtca	480
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tcggttcatt	cgtgagacc	tgtacccgat	tttacacggc	tgagattgtg	tctgctttag	660
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gcaaacaagt	ttgttgaaga	aacaagtcat	ttgcatctgt	gtggcggtgt	tgaacatgg	840
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gagatgcaat	tcattaaaga	gcaacttctc	cccagcaact	tttgggtttg	actcgtggta	1080
tcatcttcct	ccgcgaggc	tttgttaatc	accagctgc	cgtgtctcag	gtagttcagc	1140
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aagccagcgg	gatgctgcgg	gccgggcgtg	gggggtctct	tcttcctgcg	ggcccgccgt	1320
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gcacagntgc	gctcgattct	c				1461

<210> 64
 <211> 765
 <212> DNA
 <213> Homo sapiens

<400> 64	ctcaaataca	catcacaaaa	caaattttct	ctattatttg	ggctgagtca	tcactagaga	60
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	ttctccaagt	tggggcgtca	gaggggagtc	atcatgagcg	atgttaccat	tgtgaaagaa	180
	ggttgggttc	agaagagggg	agaatatata	aaaaactgga	ggccaagata	tttcctttg	240

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ttaaaccagg tttagaattt ggaatggaga aaacctttgt aaaaattatt ttaaaatgag 420
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atcttaatga aaatgactgc ggcagtcaag agtttcaa at ccagctgcc a ggggtgaaag 660
caaccctctg catctctgaa agatttcatc agtgttatct ccttataat cataacttt 720
catgtgtatc atctgagttt cttattaaat aatctcacta taaaa 765

<210> 65
<211> 968
<212> DNA
<213> Homo sapiens

<220>
<221> -
<222> (1)..(968)
<223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

<400> 65
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ttctccaagt tggggctca gaggggagtc atcatgagcg atgttaccat tgtgaaagaa 180
ggttgggttc agaagagggg agaatatata aaaaactgga ggccaagata cttcctttg 240
aagacagatg gctcattcat aggatataaa gagaaacctc aagatgtgga tttaccttat 300
cccctaaca acttttcagt ggcaaaatgc cagttatga aaacagaacg accaaagcca 360
aacacattta taatcagatg totccagtgg actactgtt tagagagaac atttcatgt 420
gatactccag aggaaaggga agaatggaca gaagctatcc aggctgtac agacagactg 480
cagaggcaag aagaggagag aatgaattgt agtccaactt cacaaattga taatatagg 540
gaggaagaga tggatgcctc tacaacccat cataaaagaa agacaatgaa tgattttgac 600
tatttggaaac tactaggtaa aggcactttt gggaaagtta tttgggtcg agagaaggca 660
agtggaaaat actatgctat gaagattctg aagaaagaag tcattattgc aaaggtaact 720
gatttattaa agttgattac taaattttt gttcagtgt gcatgtgtt gtgggctcat 780

gaatttacat gctaatttat gcaaattcca taaaacaacc naaatatgtt tgnagactac 840
tgctacagta attttgtgt attaatattt gtaatttttta aagtttcag acattcataa 900
tatttgtata ttatatacta aagctattct cttaggaaa tagaaatgtt tatgtttgca 960
tgtttggg 968

<210> 66
<211> 2410
<212> DNA
<213> Homo sapiens

<400> 66
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attnaacgc tcgcagtttc cccttgaaaa cctggaaaag ataagccagc tgctgtctaa 180
gagtgctcag tgtccactga gactacacta tctatcatca caatatgggt atgagaggtg 240
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gtaccgtat ctcaagttgg agaatctaattt gctggacaaa gatggccaca taaaattac 480
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cactccagaa tatctggcac cagaggtgtt agaagataat gactatggcc gagcagtaga 600
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ccaggaccat gagaaacttt ttgaattat attaatggaa gacattaaat ttcctcgaac 720
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caattatcat ctttgatct tttagtttt ccctcagtga aggctaaatg agatacactg	1440
attctaggta catttttaa ctttctagaa gagaaaaact aactagacta agaagattt	1500
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<213> Homo sapiens

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10

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tcgaggctgc agtgagtcat gattgtatga ctgcactcca gcctgggtga cagagcaaga 2040
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 gtccaaaagt agcccaggc tgtagcacag gcttacagt gatTTTGTGT tcagccgtga 2520
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 taaaaacgta ttttgcata tcggaaa 2608

<210> 82
 <211> 1237
 <212> DNA
 <213> Homo sapiens

<400> 82
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 cgcggccac ttcttgcacg aggtgcccgc ctgggtcatg tgccgcttct acaaagtgtat 180
 ggacgcctg gagccgcgg actggtgcca gttcgccgcc ctgatcgtgc ggcaccagac 240
 cgagctgcgg ctgtgcgagc gctccggca ggcacggcc agcgtcctgt ggccctggat 300
 caaccgcaac gcccgtgtgg cgcacccgt gcacatcctc acgcacctgc agctgctccg 360
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 gttgccatcc tcagcctcca ctttcctctc cccagctttt ccaggctccc agacccattc 540
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 cccctctccg ttttgcgtgc ccctctgtga gatttcccg ggcacccaca acttctcgaa 720
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ctcttcgtg tccaaaagta gcccagggtt gtagcacagg cttcacagtg attttgtgtt 1140
cagccgtgag tcacactaca tgcccccgtg aagctggca ttggtgacgt ccaggttgtc 1200
cttgagtaat aaaaacgtat gttgcaatct cgggaaaa 1237

<210> 83
<211> 1286
<212> DNA
<213> Homo sapiens

<220>
<221> -
<222> (1)..(1286)
<223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

<400> 83
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tgcttagac catctacata gcctggaaat aattataga gactaaaaac cagaaaatat 180
acttcttgat gaagaaggtc acatcaagtt aacagatttc ggcctaagta aagagtctat 240
tgaccatgaa aagaaggcat attcttttg tggaactgtg gagtatatgg ctccagaagt 300
agttaatcgt cgaggtcata ctcagagtgc tgactggtgg tctttgggtg tgttaatgtt 360
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gattcttaaa gccaaacttg gaatgccaca gttttgagt cctgaagcgc agagtcttt 480
acgaatgctt ttcaagcgaa atcctgcaaa cagattaggt gcaggaccag atggagttga 540
agaaattaaa agacattcat tttctcaac gatagactgg aataaaactgt atagaagaga 600
aattcatccg ccatttaaac ctgcaacggg caggcctgaa gatacattct attttgatcc 660
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gcagacagtt ggtgtacatt caattgttca gcagttacac aggaacagta ttcagttac 840
ttagtttat gaagtaaaag aagatattgg agttggctcc tactctgttt gcaagagatg 900

tatacataaa gctacaaaaca tggagttgc agtgaaggta aattttttt attaaaaatg 960
caattcatac agttcttggtt catgcgtgtc agtaccagtt aaaaattaca ctccccttgt 1020
tgttaaaagt gcctttgtt ataaaaaaagt taaatatctg gctagtgtac ttcagagatc 1080
ttaatctaga accctgtgag ctaaaggtaa ggtggttata tatctagttt tccagagca 1140
gtagcagttt acacctaag tgattttttt tctttttta cctcaagtga tttttaaagt 1200
atcttttac tctgagaagt ccccatttt tgctcanggt gtcagcaaat tcctcaaaat 1260
tgtgtgcaaa attttgtatg tttaca 1286

<210> 84
<211> 752
<212> DNA
<213> Homo sapiens

<220>
<221> -
<222> (1)..(752)
<223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

<400> 84
atgccgctgg cgtagctggc ggacccgtgg cagaagatgg ctgtggagag cccgtccgac 60
agcgctgaga atggacagca aattatggat gaacctatgg gagaggagga gattaaccca 120
caaactgaag aagtcagtat caaagaaatt gcaatcacac atcatgtaaa ggaaggacat 180
gaaaaggcag atccttccca gtttgaactt ttaaaagtat tagggcaggg atcatttgga 240
aaggtttct tagttaaaaa aatctcaggc tctgatgcta ggcagctta tgccatgaag 300
gtattgaaga aggccacact gaaagttcg aaccgagttc ggacaaaaat ggaacgtgat 360
atcttggtag aggttaatca tccttttatt gtcaagttgc attatgcttt tcaaactgaa 420
gggaagttgt atcttatttt ggattttctc aggggaggag atttgtttac acgcttatcc 480
aaagaggtga tgttcacaga agaagatgtc aaattctact tggctgaact tgcacttgct 540
ttagaccatc tacatagcct gggataatt tatagagact taaaaccaga aaagtaagga 600
atcatgctac taagttgaat acaatgtaat atgattgttt aggagattat aaaaaatcaa 660
gtggcttcat gaaactccca cagtaatgtn tagcgtgcct gtgcttcaca tctctgctaa 720
cactgttagtt tcatacttta aatnactcag tt 752

<210> 85
<211> 1826
<212> DNA

<213> Homo sapiens

<400> 85
cgagcgcggc gcccggc tgacccggc cgcaaggttt cgagccgact tgtcagccgg 60
ccaagaaaag gaagctccgt ccctcccg tcacccggct tccccacccc ttgtactcta 120
aactctgcas agggcgagcg ygcggccack gakgcgccga ggaggagcga ggcgcggcgg 180
gcagcggcgt gcccctgggg gagagggcgc cggakargag cggcggcgcg gcggcgakgg 240
cgccggcgcg gatggcagct gcttagcccg gcggcgcgg agcagccccg agctgtggct 300
ggccaggcgg tgcggctggg cgggggacgc cgccggcgtt gctgcccggc ccggagagat 360
gagcacggag gcggacgarg gcatcacttt ctctgtgcca cccttcgccc cctcgggctt 420
ctgcaccatc cccgagggcg gcatctgcag gaggggagga gcggcggcgg tggcggaggg 480
cgaggagcac cagctgccac cgccggcgc gggcagytgc tggaacgtgg agagcgcgc 540
tgccccctggc atcggttgc cggcggccac ctccctcgagc agtgcaccc gaggccgggg 600
cagctctgtt ggccccccca gccgacggac cacggggca tatgtgatca acgaagcgag 660
ccaaggccaa ctgggtggc ccgagagcga ggccctgcag agcttgcggg aggcgtgcga 720
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cgaaggacaa gttgcttcag atatgtattt cctagttggt cgaatctaca aagatatgtt 1560
tttggactct aatttacgg acactgaaag cagagaccat ggagcttctt ggttcaaaaa 1620

DNA sequence analysis

ggcatttcaa tctgagccaa cactacagtc aggaatataat tatgcgggcc tcctcctggc 1680
agctggacac cagtttgaat ctccctttga gctccggaaa gttggtaatt acaacttgat 1740
atttctacat ggaaatcaag aaactcgac ccaacttggt gcaaagacgg atctccgccc 1800
attctgacgg ctctccaggt tttgtc 1826

<210> 86
<211> 476
<212> DNA
<213> Homo sapiens

<400> 86
gccggcggtg gcgcggcggaa gacccggctg gtataacaag aggattgcct gatccagcca 60
agatgcagag cacttctaat catctgtggc ttttatctga tatttttaggc caaggagcta 120
ctgcaaatgt ctttcgtgga agacataagt ggatgttcaa atgagagaat ttgaagtgtt 180
aaaaaaaactc aatcacaaaaa atattgtcaa attatttgct attgaagagg agacaacaac 240
aagacataaa gtacttatta tggaaattttg tccatgtggg agtttataaca ctgttttaga 300
agaaccttct aatgcctatg gactaccaga atctgaattc ttaattgttt tgcgagatgt 360
ggtgggtgga atgaatcatc tacgagagaa tggtagatgt caccgtgata tcaagccagg 420
aaatatcatg cgtgcactat accattctct cgtagatgtat tcattccacc caccac 476

<210> 87
<211> 2131
<212> DNA
<213> Homo sapiens

<220>
<221> -
<222> (1)..(2131)
<223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

<400> 87
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caggatctta cttggatcca gagtgctagc tgcaaaagct tctggaaatt gtacacttaa 180
ctctgaagtt ttatctttaa tataggaagt gccgcgtatg atgctgtcct tgacagaaaat 240
gtggccatta agaagctcag cagacccttt cagaacaaaa cacatgccaa gagagcgtac 300
cgggagctgg tcctcatgaa gtgtgtgaac cataaaaaaca ttatttagttt attaaatgtc 360
ttcacaccccc agaaaacgct ggaggagttc caagatgttt acttagtaat ggaactgtatg 420

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gatgccaaact tatgtcaagt gattcagatg gaatttagacc atgagcgaat gtcttacctg	480
ctgtacccaaa tgggtgtgg cattaagcac ctccattctg ctggaattat tcacagggat	540
ttaaaaccaa gtaacattgt agtcaagtct gattgcacat tgaaaatcct ggactttgga	600
ctggccagga cagcaggcac aagcttcatg atgactccat atgtggtgac acgttattac	660
agagccccctg aggtcatcct ggggatgggc tacaaggaga acgtggatat atggctgtg	720
ggatgcatta tgggagaaat ggttcgccac aaaatcctct ttccaggaag ggactatatt	780
gaccagtgga ataaggtaat tgaacaacta ggaacaccat gtccagaatt catgaagaaa	840
ttgcaaccca cagtaagaaa ctatgtggag aatcgccca agtatgcggg actcaccttc	900
cccaaactct tcccagattc cctcttccca gcggactccg agcacaataa actcaaagcc	960
agccaagcca gggacttgtt gtcaaagatg ctatgtgattg acccagcaaa aagaatatac	1020
gtggacgacg cttacagca tccctacatc aacgtctggt atgaccacgc cgaagtggag	1080
gcgcctccac ctcagatata tgacaagcag ttggatgaaa gagaacacac aattgaagaa	1140
tggaaagaac ttatctacaa ggaagtaatg aattcagaag aaaagactaa aaatggtgta	1200
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gctactgtca gtgcctcag ggctctacca agacataatg cactgggtt ccacatggtc	1920
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gctgttttagt ttgttaattt attaaactgt atgtttata agaaaacatg taaagggggaa	2040
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2131

<210> 88
<211> 989
<212> DNA
<213> Homo sapiens

<400> 88
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ccccctgtatt tcacgcagct ctctaaattt actcagctcc aggctagtgt gagaaacacc 120
aacagcaggg ccatctcaga tcttcactat ggcaacttat gcaagaaact gttgaattag 180
acccgttcc tatagatgag aaaccataca agctgtggta tttatgagcc tccatttctt 240
atactactgc agtgaaccaa cattggatgt gaaaattgcc ttttgcagg gattcgataa 300
acaagtggat gtgtcatata ttgccaaaca ttacaacatg agcaaaaagca aagttgacaa 360
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tctaaaggcct attggctctg gggctcaggg catagttgt gccgcgtatg atgctgtcct 480
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gagagcgtac cgggagctgg tcctcatgaa gtgtgtgaac cataaaaaacg tgagtttgt 600
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caatatgcta cattcgattt cattgtcctc atggtagctt tctgcttaaa aatcacctaa 780
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atatttggtt atcttgctcc aaatttggagag cttcaagaaaa gaaacaagac aaacaaagcc 900
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tttaattttag aatttattaa tttaaaaatt 989

<210> 89
<211> 1818
<212> DNA
<213> Homo sapiens

<400> 89
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tgggagaatc gattgaacct ggaagtggag gttgcgggtga gccaagatca tcctgtcgca 120
ctccagcctg ggcaacaaga gcgaaactcc atctcaaaaaa gaaaaaaaaa gatatatatg 180
tgtgacttac aggtacaggt aaagttgctt ctgggtttct gggttggca tggtatttcc 240

tatgcagcca caggtctta ttttcttact taagtgcctc caacttccca taacacaaat	300
taaggcatga tgaacatcct ctctgtgctg aacatcctgt gtatgtcact tcagaaggct	360
gtgtgacggt ttcttagtc ttataccta ggggtggat ttctgggtca taggacagta	420
atttatattt atttcactaa gtattctctt tctctggctt ttgttacata ttacctgttt	480
gtcctccaga aaacttgcac caatttacat tcctaccaat agggtaggag agtgcacaat	540
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gaccaaagtg acactggaga attttatacg caacccatc gctcaacatg aagaacgaga	720
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aagggccatg tgaaactttc tgacttttgtt cttgcacag gactgaaaaa agcacatagg	1320
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tgtgatttgtt ggtcgattgg ggtgatcatg tatgagatgc tcatcggtaa gttgcatggt	1560
ttcagaggac ttttctgtg catccatgac agactttac attgatacca gcctctgttt	1620
caattggcag tgatctaagt gatttcccta cttgtcttcc aaagtgaatt gttttagaca	1680
gatgacacct ctttcagtaa gatgtatccc actccattct tgggcttact ggcattcgtc	1740
aattgctttg ctgatcattt tttatgtttt tctttcttctt ataccttcat cttctccatc	1800
tagaagctct ttttagtca	1818

<211> 2732
<212> DNA
<213> Homo sapiens

<220>
<221> -
<222> (1)..(2732)
<223> "n" can be any nucleotide 'a', 'c' , 'g' or 't'

<400> 90
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tagcaacaat tcttaaagag gtttggaag gcttagacta tctacacaga aacggtcaga 180
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cagattttgg ggtaagtgcg ttccctagcaa caggggggtga tgttaccga aataaaagtaa 300
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gctatgactt caaggctgac atgtggagtt ttggaaataac tgccattgaa ttagcaacag 420
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ccttagaaa attactttca ctgtgtcttc agaaagatcc ttccaaaagg cccacagcag 600
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tggatgagaa gagc当地 agaa gggaaagcag ct当地 ctca gaaaaagtc当地 cgaagagtaa 840
aagaagaaaa tccagagatt gcagttagtg ccagcaccat cccgc当地 aatacagtc当地 900
tctctgtgca cgactctc当地 ggccc当地 cccca atgctaatga agactacaga gaagttctt 960
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agattcctgca tgaagtgaag ctgattggg ttgctc当地 ggtgtc当地 tgatgtatgt 1260
cccttgc当地 gatgt caccctgatc tgtcatgccc caccgccacc cctactccct tcaaccctcc 1320
ctcttctgc ccatttccctc ccacccccc当地 actcccat tt octagcaaaa tcagaagatt 1380

0
1
2
3
4
5
6
7
8
9

gtgaagaggc	cggcttcaac	aaaatggat	aaaaaaataa	tttttaaaa	cttacaacac	1440
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gcacgactgt	tgctcatttt	tccaaaagct	attnaatatt	cttagcaatc	aatttggata	1560
tcccttaagt	gaaaagaatc	tgaaatacac	tcaggtggtc	ttatttattt	gcaacaaaag	1620
gaattttcta	tccagaagcc	tatttctcct	ttcattgttg	ttatttctgt	tataatactt	1680
taatttgtaca	tctgacaata	ctgcctctt	tatgttgat	ttagaaatta	atataactt	1740
aaaattaaga	tttatttagcc	aaacttgaat	tctagttta	aaactgactg	tgaattttat	1800
ttttcatata	tttatgcatt	acacacctt	gctataagaa	aaaaagggtt	ttgattatat	1860
gcttcttgca	gttaatctcg	ttatttaaac	aaaaagttt	gggtctgtct	ttggagtatt	1920
tgttaacttct	aaattttgaa	atgactgaat	taggaattt	gatgcttatt	cttttagtct	1980
gtttgcctaa	aaaccaattt	acaatctgac	tgtctcttgg	gagagggagg	tgccttgcaa	2040
actttcacat	taagaatgtg	cctgaggctg	ctttactctg	gaatagtctc	agatctaaaa	2100
tttcctctat	ataaggtggc	atatgttaag	tttgcttca	ttggaccgtt	tagaatgcta	2160
tgtaaaatgt	tgccattctg	tttagattgt	aactatatac	ccatctctga	tttggctctc	2220
cttaagtgtat	aggatttgg	attctaaagg	tgataaactt	gaaaatatca	gaatctgagt	2280
tttacttgaa	atttgcaga	atacccaggt	ggagtgaaaa	ttggaagggt	tttgtgcaat	2340
gactaaaagg	taaaacgctg	ttaaggttca	agaatcaata	ctttcaaccc	aagtagccct	2400
ctgcttgcact	gtatattatg	gaactagtaa	accttaggat	tttggaaatt	ggagtctaat	2460
ctttcaagga	ggtggctcc	caggatggta	ccattgctct	ttcctagcta	accctagata	2520
tggcagctct	ttaatgtact	tcaaaaagca	aatatatatt	actaaggaaa	aaaagttatt	2580
tataattgcc	ttgtcataat	tgttaaggtg	ttctagagcc	atttgcatac	aattnaatgt	2640
aatttcattc	cattctattt	tttacacaac	gattactcga	agatgactgc	aaaggtaaaa	2700
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<210> 91
<211> 1416
<212> DNA
<213> Homo sapiens

<220>
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<222> (1)..(1416)
<223> "n" may be any nucleotide 'a', 'c', 'g' or 't'

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<210> 92
<211> 434
<212> PRT
<213> Homosapiens

<400> 92

Met Pro Ala Arg Arg Leu Leu Leu Leu Thr Leu Leu Leu Pro Gly
1 5 10 15

Leu Gly Ile Phe Gly Ser Thr Ser Thr Val Thr Leu Pro Glu Thr Leu
20 25 30

Leu Phe Val Ser Thr Leu Asp Gly Ser Leu His Ala Val Ser Lys Arg
35 40 45

Thr Gly Ser Ile Lys Trp Thr Leu Lys Glu Asp Pro Val Leu Gln Val
50 55 60

Pro Thr His Val Glu Glu Pro Ala Phe Leu Pro Asp Pro Asn Asp Gly
65 70 75 80

Ser Leu Tyr Thr Leu Gly Ser Lys Asn Asn Glu Gly Leu Thr Lys Leu
85 90 95

Pro Phe Thr Ile Pro Glu Leu Val Gln Ala Ser Pro Cys Arg Ser Ser
100 105 110

Asp Gly Ile Leu Tyr Met Gly Lys Lys Gln Asp Ile Trp Tyr Val Ile
115 120 125

Asp Leu Leu Thr Gly Glu Lys Gln Gln Thr Leu Ser Ser Ala Phe Ala
130 135 140

Asp Ser Leu Cys Pro Ser Thr Ser Leu Leu Tyr Leu Gly Arg Thr Glu
145 150 155 160

Tyr Thr Ile Thr Met Tyr Asp Thr Lys Thr Arg Glu Leu Arg Trp Asn
165 170 175

Ala Thr Tyr Phe Asp Tyr Ala Ala Ser Leu Pro Glu Asp Glu Gly Asp
180 185 190

Tyr Lys Met Ser His Phe Val Ser Asn Gly Asp Gly Leu Val Val Thr
195 200 205

Val Asp Ser Glu Ser Gly Asp Val Leu Trp Ile Gln Asn Tyr Ala Ser
210 215 220

Pro Val Val Ala Phe Tyr Val Trp Gln Arg Glu Gly Leu Arg Lys Val
225 230 235 240

Met His Ile Asn Val Ala Val Glu Thr Leu Arg Tyr Leu Thr Phe Met
245 250 255

Ser Gly Glu Val Gly Arg Ile Thr Lys Trp Lys Tyr Pro Phe Pro Lys
260 265 270

Glu Thr Glu Ala Lys Ser Lys Leu Thr Pro Thr Leu Tyr Val Gly Lys
275 280 285

Tyr Ser Thr Ser Leu Tyr Ala Ser Pro Ser Met Val His Glu Gly Val
290 295 300

Ala Val Val Pro Arg Gly Ser Thr Leu Pro Leu Leu Glu Gly Pro Gln
305 310 315 320

Thr Asp Gly Val Thr Ile Gly Asp Lys Gly Glu Cys Val Ile Thr Pro
325 330 335

Ser Thr Asp Val Lys Phe Asp Pro Gly Leu Lys Ser Lys Asn Lys Leu
340 345 350

Asn Tyr Leu Arg Asn Tyr Trp Leu Leu Ile Gly His His Glu Thr Pro
355 360 365

Leu Ser Ala Ser Thr Lys Met Leu Glu Arg Phe Pro Asn Asn Leu Pro
370 375 380

Lys His Arg Glu Asn Val Ile Pro Ala Asp Ser Glu Lys Lys Ser Phe
385 390 395 400

Glu Glu Thr Leu Leu Gln Met Thr Ser Val Phe Ser Trp Ile Leu Asn
405 410 415

Leu Pro Ser Lys Glu Glu Val Phe Ala Phe Leu Arg Ile Phe Glu Lys
420 425 430

Asn Met

<210> 93
<211> 232
<212> PRT
<213> Homo sapiens

<400> 93

Met Tyr Ser Leu Gln Leu Gln Ser Val Ser Ser Ala Ile His Leu Cys
1 5 10 15

Asp Lys Lys Lys Met Glu Leu Ser Leu Asn Ile Pro Val Asn His Gly
20 25 30

Pro Gln Glu Glu Ser Cys Gly Ser Ser Gln Leu His Glu Asn Ser Gly
35 40 45

Ser Pro Glu Thr Ser Arg Ser Leu Pro Ala Pro Gln Asp Asn Asp Phe
50 55 60

Leu Ser Arg Lys Ala Gln Asp Cys Tyr Phe Met Lys Leu His His Cys
65 70 75 80

Pro Gly Asn His Ser Trp Asp Ser Thr Ile Ser Gly Ser Gln Arg Ala
85 90 95

Ala Phe Cys Asp His Lys Thr Thr Pro Cys Ser Ser Ala Ile Ile Asn
100 105 110

Pro Leu Ser Thr Ala Gly Asn Ser Glu Arg Leu Gln Pro Gly Ile Ala

115

120

125

Gln Gln Trp Ile Gln Ser Lys Arg Glu Asp Ile Val Asn Gln Met Thr
130 135 140

Glu Ala Cys Leu Asn Gln Ser Leu Asp Ala Leu Leu Ser Arg Asp Leu
145 150 155 160

Ile Met Lys Glu Asp Tyr Glu Leu Val Ser Thr Lys Pro Thr Arg Thr
165 170 175

Ser Lys Val Arg Gln Leu Leu Asp Thr Thr Asp Ile Gln Gly Glu Glu
180 185 190

Phe Ala Lys Val Ile Val Gln Lys Leu Lys Asp Asn Lys Gln Met Gly
195 200 205

Leu Gln Pro Tyr Pro Glu Ile Leu Val Val Ser Arg Ser Pro Ser Leu
210 215 220

Asn Leu Leu Gln Asn Lys Ser Met
225 230

<210> 94
<211> 209
<212> PRT
<213> Homo spaiens

<400> 94

Met Ala Asp Leu Glu Ala Val Leu Ala Asp Val Ser Tyr Leu Met Ala
1 5 10 15

Met Glu Lys Ser Lys Ala Thr Pro Ala Ala Arg Ala Ser Lys Lys Ile
20 25 30

Leu Leu Pro Glu Pro Ser Ile Arg Ser Val Met Gln Lys Tyr Leu Glu
35 40 45

Asp Arg Gly Glu Val Thr Phe Glu Lys Ile Phe Ser Gln Lys Leu Gly
50 55 60

Tyr Leu Leu Phe Arg Asp Phe Cys Leu Asn His Leu Glu Glu Ala Arg
65 70 75 80

Pro Leu Val Glu Phe Tyr Glu Glu Ile Lys Lys Tyr Glu Lys Leu Glu
85 90 95

Thr Glu Glu Glu Arg Val Ala Arg Ser Arg Glu Ile Phe Asp Ser Tyr
100 105 110

Ile Met Lys Glu Leu Leu Ala Cys Ser His Pro Phe Ser Lys Ser Ala
115 120 125

Thr Glu His Val Gln Gly His Leu Gly Lys Lys Gln Val Pro Pro Asp
130 135 140

Leu Phe Gln Pro Tyr Ile Glu Glu Ile Cys Gln Asn Leu Arg Gly Asp
145 150 155 160

Val Phe Gln Lys Phe Ile Glu Arg Val Ala Leu Ala Ala Gly Ala Ala
165 170 175

Thr Leu Pro Ala Val Pro Ser Cys Pro Asn Pro Gln His Pro Gly Ser
180 185 190

Gly Thr Thr Ala Arg His Leu Gln Val Gly Pro Tyr Trp Pro Arg Leu
195 200 205

Ala

<210> 95
<211> 454
<212> PRT
<213> Homo sapiens

<400> 95

Met Gly Leu Val Ser Ser Lys Lys Pro Asp Lys Glu Lys Pro Ile Lys
1 5 10 15

Glu Lys Asp Lys Gly Gln Trp Ser Pro Leu Lys Val Ser Ala Gln Asp
20 25 30

Lys Asp Ala Pro Pro Leu Pro Pro Leu Val Val Phe Asn His Leu Thr
35 40 45

Pro Pro Pro Pro Asp Glu His Leu Asp Glu Asp Lys His Phe Val Val
50 55 60

Ala Leu Tyr Asp Tyr Thr Ala Met Asn Asp Arg Asp Leu Gln Met Leu
65 70 75 80

Lys Gly Glu Lys Leu Gln Val Leu Lys Gly Thr Gly Asp Trp Trp Leu
85 90 95

Ala Arg Ser Leu Val Thr Gly Arg Glu Gly Tyr Val Pro Ser Asn Phe
100 105 110

Val Ala Arg Val Glu Ser Leu Glu Met Glu Arg Trp Phe Phe Arg Ser
115 120 125

Gln Gly Arg Lys Glu Ala Glu Arg Gln Leu Leu Ala Pro Ile Asn Lys
130 135 140

Ala Gly Ser Phe Leu Ile Arg Glu Ser Glu Thr Asn Lys Gly Ala Phe
145 150 155 160

Ser Leu Ser Val Lys Asp Val Thr Thr Gln Gly Glu Leu Ile Lys His
165 170 175

Tyr Lys Ile Arg Cys Leu Asp Glu Gly Gly Tyr Tyr Ile Ser Pro Arg
180 185 190

Ile Thr Phe Pro Ser Leu Gln Ala Leu Val Gln His Tyr Ser Ser Tyr
195 200 205

Tyr Lys Asn Asn Met Lys Val Ala Ile Lys Thr Leu Lys Glu Gly Thr
210 215 220

Met Ser Pro Glu Ala Phe Leu Gly Glu Ala Asn Val Met Lys Ala Leu
225 230 235 240

Gln His Glu Arg Leu Val Arg Leu Tyr Ala Val Val Thr Lys Glu Pro
245 250 255

Ile Tyr Ile Val Thr Glu Tyr Met Ala Arg Gly Cys Leu Leu Asp Phe
260 265 270

Leu Lys Thr Asp Glu Gly Ser Arg Leu Ser Leu Pro Arg Leu Ile Asp
275 280 285

Met Ser Ala Gln Ile Ala Glu Gly Met Ala Tyr Ile Glu Arg Met Asn
290 295 300

Ser Ile His Arg Asp Leu Arg Ala Ala Asn Ile Leu Val Ser Glu Ala
305 310 315 320

Leu Cys Cys Lys Ile Ala Asp Phe Gly Leu Ala Arg Ile Ile Asp Ser
325 330 335

Glu Tyr Thr Ala Gln Glu Gly Ala Lys Phe Pro Ile Lys Trp Thr Ala
340 345 350

Pro Glu Ala Ile His Phe Gly Val Phe Thr Ile Lys Ala Asp Val Trp
355 360 365

Ser Phe Gly Val Leu Leu Met Glu Val Val Thr Tyr Gly Arg Val Pro
370 375 380

Tyr Pro Gly Met Ser Asn Pro Glu Val Ile Arg Asn Leu Glu Arg Gly
385 390 395 400

Tyr Arg Met Pro Arg Pro Asp Thr Cys Pro Pro Glu Leu Tyr Arg Gly
405 410 415

Val Ile Ala Glu Cys Trp Arg Ser Arg Pro Glu Glu Arg Pro Thr Phe
420 425 430

Glu Phe Leu Gln Ser Val Leu Glu Asp Phe Tyr Thr Ala Thr Glu Arg
435 440 445

Gln Tyr Glu Leu Gln Pro
450

<210> 96
<211> 82
<212> PRT
<213> Homo sapiens

<400> 96

Met Glu Asn Phe Gln Lys Val Glu Lys Ile Gly Glu Gly Thr Tyr Gly
1 5 10 15

Val Val Tyr Lys Ala Arg Asn Lys Leu Thr Gly Glu Val Val Ala Leu
20 25 30

Lys Lys Ile Arg Leu Asp Thr Glu Thr Glu Gly Val Pro Ser Thr Ala
35 40 45

Ile Arg Glu Ile Ser Leu Leu Lys Glu Leu Asn His Pro Asn Ile Val
50 55 60

Lys Leu Leu Asp Val Ile His Thr Glu Asn Lys Asn Ile Ser Leu Lys
65 70 75 80

Glu Gly

<210> 97

<211> 118

<212> PRT

<213> Homo sapiens

<400> 97

Met Thr Arg Asp Glu Ala Leu Pro Asp Ser His Ser Ala Gln Asp Phe
1 5 10 15

Tyr Glu Asn Tyr Glu Pro Lys Glu Ile Leu Gly Arg Gly Val Ser Ser
20 25 30

Val Val Arg Arg Cys Ile His Lys Pro Thr Ser Gln Glu Tyr Ala Val
35 40 45

Lys Val Ile Asp Val Thr Gly Gly Ser Phe Ser Pro Glu Glu Val
50 55 60

Arg Glu Leu Arg Glu Ala Thr Leu Lys Glu Val Asp Ile Leu Arg Lys
65 70 75 80

Val Ser Gly His Pro Asn Ile Ser Ile Gln Leu Lys Asp Thr Tyr Glu
85 90 95

Thr Asn Thr Phe Phe Leu Val Phe Asp Leu Met Lys Arg Gly Glu
100 105 110

Leu Phe Asp Leu Pro His
115

<210> 98

<211> 167

<212> PRT

<213> Homo sapiens

<400> 98

Val Phe Leu Gly Arg Cys Arg Ser Val Lys Glu Phe Glu Lys Leu Asn
1 5 10 15

Arg Ile Gly Glu Gly Thr Tyr Gly Ile Val Tyr Arg Ala Arg Asp Thr
20 25 30

Gln Thr Asp Glu Ile Val Ala Leu Lys Lys Val Arg Met Asp Lys Glu
35 40 45

Lys Asp Gly Ile Pro Ile Ser Ser Leu Arg Glu Ile Thr Leu Leu Leu
50 55 60

Arg Leu Arg His Pro Asn Ile Val Glu Leu Lys Glu Val Val Val Gly
65 70 75 80

Asn His Leu Glu Ser Ile Phe Leu Val Met Gly Tyr Cys Glu Gln Asp
85 90 95

Leu Ala Ser Leu Leu Glu Asn Met Pro Thr Pro Phe Ser Glu Ala Gln
100 105 110

Val Lys Cys Ile Val Leu Gln Val Leu Arg Gly Leu Gln Tyr Leu His
115 120 125

Arg Asn Phe Ile Ile His Arg Asp Leu Lys Val Ser Asn Leu Leu Met
130 135 140

Thr Asp Lys Gly Cys Val Lys Thr Gly Gly Cys Asn Leu Gly Gln Ala
145 150 155 160

Trp Ser Leu Asp Gly Thr Trp
165

<210> 99
<211> 141
<212> PRT
<213> Homo sapiens

<400> 99

Met Ser Ser Ala Gly Gly Val Ser Arg Arg Leu Ala Ala Val Arg Ser
1 5 10 15

Thr Val Leu Cys Arg Ala Val Gly Cys Ile Leu Ala Glu Leu Leu Ala
20 25 30

His Arg Pro Leu Leu Pro Gly Thr Ser Glu Ile His Gln Ile Asp Leu
35 40 45

Ile Val Gln Leu Leu Gly Thr Pro Ser Glu Asn Ile Trp Pro Gly Phe
50 55 60

Ser Lys Leu Pro Leu Val Gly Gln Tyr Ser Leu Arg Lys Gln Pro Tyr
65 70 75 80

Asn Asn Leu Lys His Lys Phe Pro Trp Leu Ser Glu Ala Gly Leu Arg

85

90

95

Leu Leu His Phe Leu Phe Met Tyr Asp Pro Lys Lys Arg Ala Thr Ala
100 105 110

Gly Asp Cys Leu Glu Ser Ser Tyr Phe Lys Glu Lys Pro Leu Arg Leu
115 120 125

Pro Ile Ser Gly Val Cys Glu Gly Cys Arg Glu Pro Gly
130 135 140

<210> 100

<211> 119

<212> PRT

<213> Homo sapiens

<400> 100

Val Phe Leu Gly Arg Cys Arg Ser Val Lys Glu Phe Glu Lys Leu Asn
1 5 10 15

Arg Ile Gly Glu Gly Thr Tyr Gly Ile Val Tyr Arg Ala Arg Asp Thr
20 25 30

Gln Thr Asp Glu Ile Val Ala Leu Lys Lys Val Arg Met Asp Lys Glu
35 40 45

Lys Asp Gly Ile Pro Ile Ser Ser Leu Arg Glu Ile Thr Leu Leu Leu
50 55 60

Arg Leu Arg His Pro Asn Ile Leu Pro Ala Arg Ala Pro Trp Lys Gly
65 70 75 80

Arg Ser Gly Gly Ser Ile Arg Gly Cys Arg Gly Leu Met Trp Ser Ser
85 90 95

Ser Leu Cys Trp Lys Cys Ala Thr Thr Ala Ser Trp Glu Glu Trp Trp
100 105 110

Val Gln Ser Pro Arg Cys Leu
115

<210> 101

<211> 756

<212> PRT

<213> Homo sapiens

<400> 101

Met Gly Glu Ala Glu Lys Phe His Tyr Ile Tyr Ser Cys Asp Leu Asp
1 5 10 15

Ile Asn Val Gln Leu Lys Ile Gly Ser Leu Glu Gly Lys Arg Glu Gln
20 25 30

Lys Ser Tyr Lys Ala Val Leu Glu Asp Pro Met Leu Lys Phe Ser Gly
35 40 45

Leu Tyr Gln Glu Thr Cys Ser Asp Leu Tyr Val Thr Cys Gln Val Phe
50 55 60

Ala Glu Gly Lys Pro Leu Ala Leu Pro Val Arg Thr Ser Tyr Lys Ala
65 70 75 80

Phe Ser Thr Arg Trp Asn Trp Asn Glu Trp Leu Lys Leu Pro Val Lys
85 90 95

Tyr Pro Asp Leu Pro Arg Asn Ala Gln Val Ala Leu Thr Ile Trp Asp
100 105 110

Val Tyr Gly Pro Gly Lys Ala Val Pro Val Gly Gly Thr Thr Val Ser
115 120 125

Leu Phe Gly Lys Tyr Gly Met Phe Arg Gln Gly Met His Asp Leu Lys
130 135 140

Val Trp Pro Asn Val Glu Ala Asp Gly Ser Glu Pro Thr Lys Thr Pro
145 150 155 160

Gly Arg Thr Ser Ser Thr Leu Ser Glu Asp Gln Met Ser Arg Leu Ala
165 170 175

Lys Leu Thr Lys Ala His Arg Gln Gly His Met Val Lys Val Asp Trp
180 185 190

Leu Asp Arg Leu Thr Phe Arg Glu Ile Glu Met Ile Asn Glu Ser Val
195 200 205

Lys Arg Ser Ser Asn Phe Met Tyr Leu Met Gly Gly Phe Arg Cys Val
210 215 220

Lys Cys Asp Asp Lys Glu Tyr Gly Ile Val Tyr Tyr Glu Lys Asp Gly
225 230 235 240

Asp Glu Ser Ser Pro Ile Leu Thr Ser Phe Glu Leu Val Lys Val Pro
245 250 255

Asp Pro Gln Met Ser Leu Glu Asn Leu Val Glu Ser Lys His His Asn
260 265 270

Leu Pro Arg Ser Leu Arg Ser Gly Pro Ser Asp His Asp Leu Lys Pro
275 280 285

Tyr Pro Ser Pro Arg Asp Gln Leu Lys Asn Ile Val Ser Tyr Pro Pro
290 295 300

Ser Lys Pro Pro Thr Tyr Glu Glu Gln Asp Leu Val Trp Glu Phe Arg
305 310 315 320

Tyr Tyr Leu Thr Asn Gln Asp Lys Ala Leu Thr Lys Ile Leu Thr Ser
325 330 335

Val Ile Trp Asp Leu Pro Gln Glu Ala Lys Gln Ala Leu Ala Leu Leu
340 345 350

Gly Lys Trp Asn Pro Met Asp Val Glu Asp Ser Leu Glu Leu Ile Ser
355 360 365

Ser His Tyr Thr Asn Pro Thr Val Arg Arg Tyr Ala Val Ala Arg Leu
370 375 380

Arg Gln Ala Asp Asp Glu Asp Leu Leu Met Tyr Leu Leu Gln Leu Val
385 390 395 400

Gln Ala Leu Lys Tyr Glu Asn Phe Asp Asp Ile Lys Asn Gly Leu Glu
405 410 415

Pro Thr Lys Lys Asp Ser Gln Ser Ser Val Ser Glu Asn Val Ser Asn
420 425 430

Ser Gly Ile Asn Ser Ala Glu Ile Asp Ser Ser Gln Ile Ile Thr Ser
435 440 445

Pro Leu Pro Ser Val Ser Ser Pro Pro Pro Ala Ser Lys Thr Lys Glu
450 455 460

Val Pro Asp Gly Glu Asn Leu Glu Gln Asp Leu Cys Thr Phe Leu Ile
465 470 475 480

Ser Arg Ala Cys Lys Asn Ser Thr Leu Ala Asn Tyr Leu Tyr Trp Tyr
485 490 495

Val Ile Val Glu Cys Glu Asp Gln Asp Thr Gln Gln Arg Asp Pro Lys
500 505 510

Thr His Glu Met Tyr Leu Asn Val Met Arg Arg Phe Ser Gln Ala Leu
515 520 525

Leu Lys Gly Asp Lys Ser Val Arg Val Met Arg Ser Leu Leu Ala Ala
530 535 540

Gln Gln Thr Phe Val Asp Arg Leu Val His Leu Met Lys Ala Val Gln
545 550 555 560

Arg Glu Ser Gly Asn Arg Lys Lys Asn Glu Arg Leu Gln Ala Leu
565 570 575

Leu Gly Asp Asn Glu Lys Met Asn Leu Ser Asp Val Glu Leu Ile Pro
580 585 590

Leu Pro Leu Glu Pro Gln Val Lys Ile Arg Gly Ile Ile Pro Glu Thr
595 600 605

Ala Thr Leu Phe Lys Ser Ala Leu Met Pro Ala Gln Leu Phe Phe Lys
610 615 620

Thr Glu Asp Gly Gly Lys Tyr Pro Val Ile Phe Lys His Gly Asp Asp
625 630 635 640

Leu Arg Gln Asp Gln Leu Ile Leu Gln Ile Ile Ser Leu Met Asp Lys
645 650 655

Leu Leu Arg Lys Glu Asn Leu Asp Leu Lys Leu Thr Pro Tyr Lys Val
660 665 670

Leu Ala Thr Ser Thr Lys His Gly Phe Met Gln Phe Ile Gln Ser Val
675 680 685

Pro Val Ala Glu Val Leu Asp Thr Glu Gly Ser Ile Gln Asn Phe Phe
690 695 700

Arg Lys Tyr Ala Pro Ser Glu Asn Gly Pro Asn Gly Ile Ser Ala Glu
705 710 715 720

Val Met Asp Thr Tyr Val Lys Ser Cys Ala Gly Tyr Cys Val Ile Thr
725 730 735

Tyr Ile Leu Gly Val Gly Asp Arg His Leu Asp Asn Leu Leu Thr
740 745 750

Lys Thr Gly Gly
755

<210> 102

<211> 508

<212> PRT

<213> Homo sapiens

<400> 102

Met Gly Glu Ala Glu Lys Phe His Tyr Ile Tyr Ser Cys Asp Leu Asp
1 5 10 15

Ile Asn Val Gln Leu Lys Ile Gly Ser Leu Glu Gly Lys Arg Glu Gln
20 25 30

Lys Ser Tyr Lys Ala Val Leu Glu Asp Pro Met Leu Lys Phe Ser Gly
35 40 45

Leu Tyr Gln Glu Thr Cys Ser Asp Leu Tyr Val Thr Cys Gln Val Phe
50 55 60

Ala Glu Gly Lys Pro Leu Ala Leu Pro Val Arg Thr Ser Tyr Lys Ala
65 70 75 80

Phe Ser Thr Arg Trp Asn Trp Asn Glu Trp Leu Lys Leu Pro Val Lys
85 90 95

Tyr Pro Asp Leu Pro Arg Asn Ala Gln Val Ala Leu Thr Ile Trp Asp
100 105 110

Val Tyr Gly Pro Gly Lys Ala Val Pro Val Gly Gly Thr Thr Val Ser
115 120 125

Leu Phe Gly Lys Tyr Gly Met Phe Arg Gln Gly Met His Asp Leu Lys
130 135 140

Val Trp Pro Asn Val Glu Ala Asp Gly Ser Glu Pro Thr Lys Thr Pro

145 150 155 160
Gly Arg Thr Ser Ser Thr Leu Ser Glu Asp Gln Met Ser Arg Leu Ala
165 170 175
Lys Leu Thr Lys Ala His Arg Gln Gly His Met Val Lys Val Asp Trp
180 185 190
Leu Asp Arg Leu Thr Phe Arg Glu Ile Glu Met Ile Asn Glu Ser Val
195 200 205
Lys Arg Ser Ser Asn Phe Met Tyr Leu Met Gly Gly Phe Arg Cys Val
210 215 220
Lys Cys Asp Asp Lys Glu Tyr Gly Ile Val Tyr Tyr Glu Lys Asp Gly
225 230 235 240
Asp Glu Ser Ser Pro Ile Leu Thr Ser Phe Glu Leu Val Lys Val Pro
245 250 255
Asp Pro Gln Met Ser Leu Glu Asn Leu Val Glu Ser Lys His His Asn
260 265 270
Leu Pro Arg Ser Leu Arg Ser Gly Pro Ser Asp His Asp Leu Lys Pro
275 280 285
Tyr Pro Ser Pro Arg Asp Gln Leu Lys Asn Ile Val Ser Tyr Pro Pro
290 295 300
Ser Lys Pro Pro Thr Tyr Glu Glu Gln Asp Leu Val Trp Glu Phe Arg
305 310 315 320
Tyr Tyr Leu Thr Asn Gln Asp Lys Ala Leu Thr Lys Ile Leu Thr Ser
325 330 335
Val Ile Trp Asp Leu Pro Gln Glu Ala Lys Gln Ala Leu Ala Leu Leu
340 345 350
Gly Lys Trp Asn Pro Met Asp Val Glu Asp Ser Leu Glu Leu Ile Ser
355 360 365
Ser His Tyr Thr Asn Pro Thr Val Arg Arg Tyr Ala Val Ala Arg Leu
370 375 380
Arg Gln Ala Asp Asp Glu Asp Leu Leu Met Tyr Leu Leu Gln Leu Val
385 390 395 400
Gln Ala Leu Lys Tyr Glu Asn Phe Asp Asp Ile Lys Asn Gly Leu Glu
405 410 415
Pro Thr Lys Lys Asp Ser Gln Ser Ser Val Ser Glu Asn Val Ser Asn
420 425 430
Ser Gly Ile Asn Ser Ala Glu Ile Asp Ser Ser Gln Ile Ile Thr Ser
435 440 445
Pro Leu Pro Ser Val Ser Ser Pro Pro Pro Ala Ser Lys Thr Lys Glu

450

455

460

Val Pro Asp Gly Glu Asn Leu Glu Gln Asp Leu Cys Thr Phe Leu Ile
465 470 475 480

Ser Arg Ala Cys Lys Asn Ser Thr Leu Ala Asn Tyr Leu Tyr Trp Tyr
485 490 495

Val Lys Ile Ile Phe Cys Leu Phe Ser Tyr Tyr Pro
500 505

<210> 103

<211> 140

<212> PRT

<213> Homo sapiens

<400> 103

Met Gly Asn Ala Ala Ala Lys Lys Gly Ser Glu Gln Glu Ser Val
1 5 10 15

Lys Glu Phe Leu Ala Lys Ala Lys Glu Asp Phe Leu Lys Lys Trp Glu
20 25 30

Ser Pro Ala Gln Asn Thr Ala His Leu Asp Gln Phe Glu Arg Ile Lys
35 40 45

Thr Leu Gly Thr Gly Ser Phe Gly Arg Val Met Leu Val Lys His Lys
50 55 60

Glu Thr Gly Asn His Tyr Ala Met Lys Ile Leu Asp Lys Gln Lys Val
65 70 75 80

Val Lys Leu Lys Gln Ile Glu His Thr Leu Asn Glu Lys Arg Ile Leu
85 90 95

Gln Ala Val Asn Phe Pro Phe Leu Val Lys Leu Glu Phe Ser Phe Lys
100 105 110

Asp Asn Ser Asn Leu Tyr Met Val Met Glu Tyr Val Pro Gly Gly Glu
115 120 125

Met Phe Ser His Leu Arg Arg Ile Gly Arg Phe Arg
130 135 140

<210> 104

<211> 156

<212> PRT

<213> Homo sapiens

<400> 104

Met Val Val Phe Asn Gly Leu Leu Lys Ile Lys Ile Cys Glu Ala Val
1 5 10 15

Ser Leu Lys Pro Thr Ala Trp Ser Leu Arg His Ala Val Gly Pro Arg
20 25 30

Pro Gln Thr Phe Leu Leu Asp Pro Tyr Ile Ala Leu Asn Val Asp Asp
35 40 45

Ser Arg Ile Gly Gln Thr Ala Thr Lys Gln Lys Thr Asn Ser Pro Ala
50 55 60

Trp His Asp Glu Phe Val Thr Asp Val Cys Asn Gly Arg Lys Ile Glu
65 70 75 80

Leu Ala Val Phe His Asp Ala Pro Ile Gly Tyr Asp Asp Phe Val Ala
85 90 95

Asn Cys Thr Ile Gln Phe Glu Glu Leu Leu Gln Asn Gly Ser Arg His
100 105 110

Phe Glu Asp Trp Ile Asp Leu Glu Pro Glu Gly Arg Val Tyr Val Ile
115 120 125

Ile Asp Leu Ser Gly Ser Ser Gly Glu Val Lys Ile Pro Asn Ser Ala
130 135 140

Phe Cys Glu Arg Glu Arg Val Glu Met Arg His Ser
145 150 155

<210> 105
<211> 520
<212> PRT
<213> Homo sapiens

<400> 105

Met Ile Leu Ile Pro Arg Met Leu Leu Val Leu Phe Leu Leu Pro
1 5 10 15

Ile Leu Ser Ser Ala Lys Ala Gln Val Asn Pro Ala Ile Cys Arg Tyr
20 25 30

Pro Leu Gly Met Ser Gly Gly Gln Ile Pro Asp Glu Asp Ile Thr Ala
35 40 45

Ser Ser Gln Trp Ser Glu Ser Thr Ala Ala Lys Tyr Gly Arg Leu Asp
50 55 60

Ser Glu Glu Gly Asp Gly Ala Trp Cys Pro Glu Ile Pro Val Glu Pro
65 70 75 80

Asp Asp Leu Lys Glu Phe Leu Gln Ile Asp Leu His Thr Leu His Phe
85 90 95

Ile Thr Leu Val Gly Thr Gln Gly Arg His Ala Gly Gly His Gly Ile
100 105 110

Glu Phe Ala Pro Met Tyr Lys Ile Asn Tyr Ser Arg Asp Gly Thr Arg
115 120 125

Trp Ile Ser Trp Arg Asn Arg His Gly Lys Gln Val Leu Asp Gly Asn

130 135 140

Ser Asn Pro Tyr Asp Ile Phe Leu Lys Asp Leu Glu Pro Pro Ile Val
145 150 155 160

Ala Arg Phe Val Arg Phe Ile Pro Val Thr Asp His Ser Met Asn Val
165 170 175

Cys Met Arg Val Glu Leu Tyr Gly Cys Val Trp Leu Asp Gly Leu Val
180 185 190

Ser Tyr Asn Ala Pro Ala Gly Gln Gln Phe Val Leu Pro Gly Gly Ser
195 200 205

Ile Ile Tyr Leu Asn Asp Ser Val Tyr Asp Gly Ala Val Gly Tyr Ser
210 215 220

Met Thr Glu Gly Leu Gly Gln Leu Thr Asp Gly Val Ser Gly Leu Asp
225 230 235 240

Asp Phe Thr Gln Thr His Glu Tyr His Val Trp Pro Gly Tyr Asp Tyr
245 250 255

Val Gly Trp Arg Asn Glu Ser Ala Thr Asn Gly Tyr Ile Glu Ile Met
260 265 270

Phe Glu Phe Asp Arg Ile Arg Asn Phe Thr Thr Met Lys Val His Cys
275 280 285

Asn Asn Met Phe Ala Lys Gly Val Lys Ile Phe Lys Glu Val Gln Cys
290 295 300

Tyr Phe Arg Ser Glu Ala Ser Glu Trp Glu Pro Asn Ala Ile Ser Phe
305 310 315 320

Pro Leu Val Leu Asp Asp Val Asn Pro Ser Ala Arg Phe Val Thr Val
325 330 335

Pro Leu His His Arg Met Ala Ser Ala Ile Lys Cys Gln Tyr His Phe
340 345 350

Ala Asp Thr Trp Met Met Phe Ser Glu Ile Thr Phe Gln Ser Asp Ala
355 360 365

Ala Met Tyr Asn Asn Ser Glu Ala Leu Pro Thr Ser Pro Met Ala Pro
370 375 380

Thr Thr Tyr Asp Pro Met Leu Lys Val Asp Asp Ser Asn Thr Arg Ile
385 390 395 400

Leu Ile Gly Cys Leu Val Ala Ile Ile Phe Ile Leu Leu Ala Ile Ile
405 410 415

Val Ile Ile Leu Trp Arg Gln Phe Trp Gln Lys Met Leu Glu Lys Ala
420 425 430

Ser Arg Arg Met Leu Asp Asp Glu Met Thr Val Ser Leu Ser Leu Pro

435

440

445

Ser Asp Ser Ser Met Phe Asn Asn Asn Arg Ser Ser Ser Pro Ser Glu
450 455 460

Gln Gly Ser Asn Ser Thr Tyr Asp Arg Ile Phe Pro Leu Arg Pro Asp
465 470 475 480

Tyr Gln Glu Pro Ser Arg Leu Ile Arg Lys Leu Pro Glu Phe Ala Pro
485 490 495

Gly Glu Glu Glu Ser Gly Glu Asp Asp Val Val Glu Gln Gly Val Lys
500 505 510

Gly Glu Thr Ser Ala Ser Ile Arg
515 520

<210> 106

<211> 284

<212> PRT

<213> Homo sapiens

<400> 106

Met Ala Asn Phe Gln Glu His Leu Ser Cys Ser Ser Ser Pro His Leu
1 5 10 15

Pro Phe Ser Glu Ser Lys Thr Phe Asn Gly Leu Gln Asp Glu Leu Thr
20 25 30

Ala Met Gly Asn His Pro Ser Pro Lys Leu Leu Glu Asp Gln Gln Glu
35 40 45

Lys Gly Met Val Arg Thr Glu Leu Ile Glu Ser Val His Ser Pro Val
50 55 60

Thr Thr Thr Val Leu Thr Ser Val Ser Glu Asp Ser Arg Asp Gln Phe
65 70 75 80

Glu Asn Ser Val Leu Gln Leu Arg Glu His Asp Glu Ser Glu Thr Ala
85 90 95

Val Ser Gln Gly Asn Ser Asn Thr Val Asp Gly Glu Ser Thr Ser Gly
100 105 110

Thr Glu Asp Ile Lys Ile Gln Phe Ser Arg Ser Gly Ser Gly
115 120 125

Gly Phe Leu Glu Gly Leu Phe Gly Cys Leu Arg Pro Val Trp Asn Ile
130 135 140

Ile Gly Lys Ala Tyr Ser Thr Asp Tyr Lys Phe Met Gln Gln Asp Thr
145 150 155 160

Trp Glu Val Pro Phe Glu Glu Ile Ser Glu Leu Gln Trp Leu Gly Ser
165 170 175

Gly Ala Gln Gly Ala Val Phe Leu Gly Lys Phe Arg Ala Glu Glu Val
180 185 190

Ala Ile Lys Lys Val Arg Glu Gln Asn Glu Thr Asp Ile Lys His Leu
195 200 205

Arg Lys Leu Lys His Pro Asn Ile Ile Ala Phe Lys Gly Val Cys Thr
210 215 220

Gln Ala Pro Cys Tyr Cys Ile Ile Met Glu Tyr Cys Ala His Gly Gln
225 230 235 240

Leu Tyr Glu Val Leu Arg Ala Gly Arg Lys Ile Thr Pro Arg Leu Leu
245 250 255

Val Asp Trp Ser Thr Gly Ile Ala Ser Gly Met Asn Tyr Leu His Leu
260 265 270

His Lys Ile Ile His Arg Asp Leu Lys Ser Pro Lys
275 280

<210> 107
<211> 185
<212> PRT
<213> Homo sapiens

<400> 107

Met Cys Gly Gln Arg Trp Ile His Asn Phe Thr Cys Leu Ala Phe Leu
1 5 10 15

Phe His Thr Leu Lys Ser Gly Asn Lys Ser Val His Leu Arg Lys Ala
20 25 30

Ser Ser Pro Asn Leu His Arg Arg Gln Trp Glu Lys Asn Val Pro Asn
35 40 45

Thr Ala Leu Thr Ala Leu Glu Asn Ala Ser Ile Leu Thr Ser Ser Leu
50 55 60

Thr Ala Glu Asp Asp Arg Gly Gly Ser Val Ile Lys Tyr Ser Lys Asn
65 70 75 80

Thr Thr Arg Lys Gln Trp Leu Lys Glu Thr Pro Asp Thr Leu Leu Asn
85 90 95

Ile Leu Lys Asn Ala Asp Leu Ser Leu Ala Phe Gln Thr Tyr Thr Ile
100 105 110

Tyr Arg Pro Gly Ser Glu Gly Phe Leu Lys Gly Pro Leu Ser Glu Glu
115 120 125

Thr Glu Ala Ser Asp Ser Val Asp Gly Gly His Asp Ser Val Ile Leu
130 135 140

Asp Pro Glu Arg Leu Glu Pro Gly Leu Asp Glu Glu Asp Thr Asp Phe
145 150 155 160

Glu Glu Glu Asp Asp Asn Pro Asp Trp Val Ser Glu Leu Lys Lys Arg
165 170 175

Ala Gly Trp Gln Gly Leu Cys Asp Arg
180 185

<210> 108
<211> 83
<212> PRT
<213> Homo sapiens

<400> 108

Met Ala Pro Pro Ser Glu Glu Thr Pro Leu Ile Pro Gln Arg Ser Cys
1 5 10 15

Ser Leu Leu Ser Thr Glu Ala Gly Ala Leu His Val Leu Leu Pro Ala
20 25 30

Arg Gly Pro Gly Pro Pro Gln Arg Leu Ser Phe Ser Phe Gly Val Pro
35 40 45

Val Arg Pro Val Gly Ala Asn Gly Pro Pro Leu Thr Ser Gly Phe Leu
50 55 60

Gly Gly Trp Ala Glu Ala Ser Val Gln Arg Gly Leu Trp Lys Cys Leu
65 70 75 80

Leu Thr Glu

<210> 109
<211> 213
<212> PRT
<213> Homo sapiens

<400> 109

Met Ala Glu Ser Ala Gly Ala Ser Ser Phe Phe Pro Leu Val Val Leu
1 5 10 15

Leu Leu Ala Gly Ser Gly Ser Gly Pro Arg Gly Val Gln Ala Leu
20 25 30

Leu Cys Ala Cys Thr Ser Cys Leu Gln Ala Asn Tyr Thr Cys Glu Thr
35 40 45

Asp Gly Ala Cys Met Val Ser Ile Phe Asn Leu Asp Gly Met Glu His
50 55 60

His Val Arg Thr Cys Ile Pro Lys Val Glu Leu Val Pro Ala Gly Lys
65 70 75 80

Pro Phe Tyr Cys Leu Ser Ser Glu Asp Leu Arg Asn Thr His Cys Cys
85 90 95

Tyr Thr Asp Tyr Cys Asn Arg Ile Asp Leu Arg Val Pro Ser Gly His
100 105 110

Leu Lys Glu Pro Glu His Pro Ser Met Trp Gly Pro Val Glu Leu Val
115 120 125

Gly Ile Ile Ala Gly Pro Val Phe Leu Leu Phe Leu Ile Ile Ile Ile
130 135 140

Val Phe Leu Val Ile Asn Tyr His Gln Arg Val Tyr His Asn Arg Gln
145 150 155 160

Arg Leu Asp Met Glu Asp Pro Ser Cys Glu Met Cys Leu Ser Lys Asp
165 170 175

Lys Thr Leu Gln Asp Leu Val Tyr Asp Leu Ser Thr Ser Gly Ser Gly
180 185 190

Ser Gly Thr Lys Phe Phe Arg Ala Ser Cys Leu Trp Leu Ala Phe Ile
195 200 205

Ser Phe Pro Ala Gly
210

<210> 110
<211> 383
<212> PRT
<213> Homo sapiens

<400> 110

Met Asp Glu Gln Glu Ala Leu Asn Ser Ile Met Asn Asp Leu Val Ala
1 5 10 15

Leu Gln Met Asn Arg Arg His Arg Met Pro Gly Tyr Glu Thr Met Lys
20 25 30

Asn Lys Asp Thr Gly His Ser Asn Arg Gln Ser Asp Val Arg Ile Lys
35 40 45

Phe Glu His Asn Gly Glu Arg Arg Ile Ile Ala Phe Ser Arg Pro Val
50 55 60

Lys Tyr Glu Asp Val Glu His Lys Val Thr Thr Val Phe Gly Gln Pro
65 70 75 80

Leu Asp Leu His Tyr Met Asn Asn Glu Leu Ser Ile Leu Leu Lys Asn
85 90 95

Gln Asp Asp Leu Asp Lys Ala Ile Asp Ile Leu Asp Arg Ser Ser Ser
100 105 110

Met Lys Ser Leu Arg Ile Leu Leu Ser Gln Asp Arg Asn His Asn
115 120 125

Ser Ser Ser Pro His Ser Glu Val Ser Arg Gln Val Arg Ile Lys Ala
130 135 140

Ser Gln Ser Ala Gly Asp Ile Asn Thr Ile Tyr Gln Pro Pro Glu Pro
145 150 155 160

Arg Ser Arg His Leu Ser Val Ser Ser Gln Asn Pro Gly Arg Ser Ser
165 170 175

Pro Pro Pro Gly Tyr Val Pro Glu Arg Gln Gln His Ile Ala Arg Gln
180 185 190

Gly Ser Tyr Thr Ser Ile Asn Ser Glu Gly Glu Phe Ile Pro Glu Thr
195 200 205

Ser Glu Gln Cys Met Leu Asp Pro Leu Ser Ser Ala Glu Asn Ser Leu
210 215 220

Ser Gly Ser Cys Gln Ser Leu Asp Arg Ser Ala Asp Ser Pro Ser Phe
225 230 235 240

Arg Lys Ser Arg Met Ser Arg Ala Gln Ser Phe Pro Asp Asn Arg Gln
245 250 255

Glu Tyr Ser Asp Arg Glu Thr Gln Leu Tyr Asp Lys Gly Val Lys Gly
260 265 270

Gly Thr Tyr Pro Arg Arg Tyr His Val Ser Val His His Lys Asp Tyr
275 280 285

Ser Asp Gly Arg Arg Thr Phe Pro Arg Ile Arg Arg His Gln Gly Asn
290 295 300

Leu Phe Thr Leu Val Pro Ser Ser Arg Ser Leu Ser Thr Asn Gly Glu
305 310 315 320

Asn Met Gly Leu Ala Val Gln Tyr Leu Asp Pro Arg Gly Arg Leu Arg
325 330 335

Ser Ala Asp Ser Glu Asn Ala Leu Ser Val Gln Glu Arg Asn Val Pro
340 345 350

Thr Lys Cys Glu Glu Leu Ser Leu Ala Arg Arg Arg Leu Pro Arg Trp
355 360 365

Ser Gln Thr Ser Tyr Gly Gly Lys Gln Leu Gly Pro Trp Asp Pro
370 375 380

<210> 111
<211> 414
<212> PRT
<213> Homo sapiens

<400> 111

Met Asp Glu Gln Glu Ala Leu Asn Ser Ile Met Asn Asp Leu Val Ala
1 5 10 15

Leu Gln Met Asn Arg Arg His Arg Met Pro Gly Tyr Glu Thr Met Lys

20

25

30

Asn Lys Asp Thr Gly His Ser Asn Arg Gln Lys Lys His Asn Ser Ser
35 40 45

Ser Ser Ala Leu Leu Asn Ser Pro Thr Val Thr Thr Ser Ser Cys Ala
50 55 60

Gly Ala Ser Glu Lys Lys Phe Leu Ser Asp Val Arg Ile Lys Phe
65 70 75 80

Glu His Asn Gly Glu Arg Arg Ile Ile Ala Phe Ser Arg Pro Val Lys
85 90 95

Tyr Glu Asp Val Glu His Lys Val Thr Thr Val Phe Gly Gln Pro Leu
100 105 110

Asp Leu His Tyr Met Asn Asn Glu Leu Ser Ile Leu Leu Lys Asn Gln
115 120 125

Asp Asp Leu Asp Lys Ala Ile Asp Ile Leu Asp Arg Ser Ser Ser Met
130 135 140

Lys Ser Leu Arg Ile Leu Leu Ser Gln Asp Arg Asn His Asn Ser
145 150 155 160

Ser Ser Pro His Ser Glu Val Ser Arg Gln Val Arg Ile Lys Ala Ser
165 170 175

Gln Ser Ala Gly Asp Ile Asn Thr Ile Tyr Gln Pro Pro Glu Pro Arg
180 185 190

Ser Arg His Leu Ser Val Ser Ser Gln Asn Pro Gly Arg Ser Ser Pro
195 200 205

Pro Pro Gly Tyr Val Pro Glu Arg Gln Gln His Ile Ala Arg Gln Gly
210 215 220

Ser Tyr Thr Ser Ile Asn Ser Glu Gly Glu Phe Ile Pro Glu Thr Ser
225 230 235 240

Glu Gln Cys Met Leu Asp Pro Leu Ser Ser Ala Glu Asn Ser Leu Ser
245 250 255

Gly Ser Cys Gln Ser Leu Asp Arg Ser Ala Asp Ser Pro Ser Phe Arg
260 265 270

Lys Ser Arg Met Ser Arg Ala Gln Ser Phe Pro Asp Asn Arg Gln Glu
275 280 285

Tyr Ser Asp Arg Glu Thr Gln Leu Tyr Asp Lys Gly Val Lys Gly Gly
290 295 300

Thr Tyr Pro Arg Arg Tyr His Val Ser Val His His Lys Asp Tyr Ser
305 310 315 320

Asp Gly Arg Arg Thr Phe Pro Arg Ile Arg Arg His Gln Gly Asn Leu

325

330

335

Phe Thr Leu Val Pro Ser Ser Arg Ser Leu Ser Thr Asn Gly Glu Asn
340 345 350

Met Gly Leu Ala Val Gln Tyr Leu Asp Pro Arg Gly Arg Leu Arg Ser
355 360 365

Ala Asp Ser Glu Asn Ala Leu Ser Val Gln Glu Arg Asn Val Pro Thr
370 375 380

Lys Cys Glu Glu Leu Ser Leu Ala Arg Arg Arg Leu Pro Arg Trp Ser
385 390 395 400

Gln Thr Ser Tyr Gly Gly Lys Gln Leu Gly Pro Trp Asp Pro
405 410

<210> 112

<211> 201

<212> PRT

<213> Homo sapiens

<400> 112

Met Ala Lys Gln Tyr Asp Ser Val Glu Cys Pro Phe Cys Asp Glu Val
1 5 10 15

Ser Lys Tyr Glu Lys Leu Ala Lys Ile Gly Gln Gly Thr Phe Gly Glu
20 25 30

Val Phe Lys Ala Arg His Arg Lys Thr Gly Gln Lys Val Ala Leu Lys
35 40 45

Lys Val Leu Met Glu Asn Glu Lys Glu Gly Phe Pro Ile Thr Ala Leu
50 55 60

Arg Glu Ile Lys Ile Leu Gln Leu Leu Lys His Glu Asn Val Val Asn
65 70 75 80

Leu Ile Glu Ile Cys Arg Thr Lys Ala Ser Pro Tyr Asn Arg Cys Lys
85 90 95

Gly Ser Ile Tyr Leu Val Phe Asp Phe Cys Glu His Asp Leu Ala Gly
100 105 110

Leu Leu Ser Asn Val Leu Val Lys Phe Thr Leu Ser Glu Ile Lys Arg
115 120 125

Val Met Gln Met Leu Leu Asn Gly Leu Tyr Tyr Asn His Asp Phe Phe
130 135 140

Trp Ser Asp Pro Met Pro Ser Asp Leu Lys Gly Met Leu Ser Thr His
145 150 155 160

Leu Thr Ser Met Phe Glu Tyr Leu Ala Pro Pro Arg Arg Lys Gly Ser
165 170 175

Gln Ile Thr Gln Gln Ser Thr Asn Gln Ser Arg Asn Pro Ala Thr Thr
180 185 190

Asn Gln Thr Glu Phe Glu Arg Val Phe
195 200

<210> 113
<211> 125
<212> PRT
<213> Homo sapiens

<400> 113

Met Ala Thr Ser Arg Tyr Glu Pro Val Ala Glu Ile Gly Val Gly Ala
1 5 10 15

Tyr Gly Thr Val Tyr Lys Ala Arg Asp Pro His Ser Gly His Phe Cys
20 25 30

Ala Leu Lys Ser Val Arg Val Pro Asn Gly Gly Gly Gly Gly Gly
35 40 45

Leu Pro Ile Ser Thr Val Arg Glu Val Ala Leu Leu Arg Arg Leu Glu
50 55 60

Ala Phe Glu His Pro Asn Val Val Arg Leu Met Asp Val Cys Ala Thr
65 70 75 80

Ser Arg Thr Asp Arg Glu Ile Lys Val Thr Leu Val Phe Glu His Val
85 90 95

Asp Gln Asp Leu Arg Thr Tyr Leu Asp Lys Ala Pro Pro Pro Gly Leu
100 105 110

Pro Ala Glu Thr Ile Lys Val Ser Gly Val Gly Arg His
115 120 125

<210> 114
<211> 45
<212> PRT
<213> Homo sapiens

<400> 114

Met Ala Thr Ser Arg Tyr Glu Pro Val Ala Glu Ile Gly Val Gly Ala
1 5 10 15

Tyr Gly Thr Val Tyr Lys Ala Arg Asp Pro His Ser Gly His Phe Cys
20 25 30

Ala Leu Lys Ser Val Arg Val Pro Thr His Leu Ser Phe
35 40 45

<210> 115
<211> 160
<212> PRT
<213> Homo sapiens

<400> 115

Met Gly Val Cys Pro Gly Lys Thr Pro Phe Cys Ser Pro Lys Pro Gln
1 5 10 15

Gly Leu Ala Arg Gly His Trp Ser Arg Arg Arg Asp Ile Cys Val Thr
20 25 30

Gly Pro Leu Pro Leu Glu Pro Arg Ala Val Tyr Cys Lys Asp Val Leu
35 40 45

Asp Ile Glu Gln Phe Ser Thr Val Lys Gly Val Asn Leu Asp His Thr
50 55 60

Asp Asp Asp Phe Tyr Ser Lys Phe Ser Thr Gly Ser Val Ser Ile Pro
65 70 75 80

Trp Gln Asn Glu Met Ile Glu Thr Glu Cys Phe Lys Glu Leu Asn Val
85 90 95

Phe Gly Pro Asn Gly Thr Leu Pro Pro Asp Leu Asn Arg Asn His Pro
100 105 110

Pro Glu Pro Pro Lys Lys Gly Leu Leu Gln Arg Leu Phe Lys Arg Gln
115 120 125

His Gln Asn Asn Ser Lys Ser Ser Pro Ser Ser Lys Thr Ser Phe Asn
130 135 140

His His Ile Asn Ser Asn His Val Ser Ser Asn Ser Thr Gly Ser Ser
145 150 155 160

<210> 116

<211> 300

<212> PRT

<213> Homo sapiens

<220>

<221> -

<222> (1)..(300)

<223> "XAA" can be any amino acid

<400> 116

Met Pro Arg Ala Arg Met Pro Xaa Pro Arg Ala His Ser Lys Ala Gly
1 5 10 15

Cys Pro Cys Gly Cys Pro Arg Asp Pro Leu Thr Leu Leu Ser Pro Ser
20 25 30

Gly His Ile Arg Ile Ser Asp Leu Gly Leu Ala Val Lys Ile Pro Glu
35 40 45

Gly Asp Leu Ile Arg Gly Arg Val Gly Thr Val Gly Tyr Met Ala Pro
50 55 60

Glu Val Leu Asn Asn Gln Arg Tyr Gly Leu Ser Pro Asp Tyr Trp Gly
65 70 75 80

Leu Gly Cys Leu Ile Tyr Glu Met Ile Glu Gly Gln Ser Pro Phe Arg
85 90 95

Gly Arg Lys Glu Lys Val Lys Arg Glu Glu Val Asp Arg Arg Val Leu
100 105 110

Glu Thr Glu Glu Val Tyr Ser His Lys Phe Ser Glu Glu Ala Lys Ser
115 120 125

Ile Cys Lys Met Leu Leu Thr Lys Asp Ala Lys Gln Arg Leu Gly Cys
130 135 140

Gln Glu Glu Gly Ala Ala Glu Val Lys Arg His Pro Phe Phe Arg Asn
145 150 155 160

Met Asn Phe Lys Arg Leu Glu Ala Gly Met Leu Asp Pro Pro Phe Val
165 170 175

Pro Asp Pro Arg Ala Val Tyr Cys Lys Asp Val Leu Asp Ile Glu Gln
180 185 190

Phe Ser Thr Val Lys Gly Val Asn Leu Asp His Thr Asp Asp Asp Phe
195 200 205

Tyr Ser Lys Phe Ser Thr Gly Ser Val Ser Ile Pro Trp Gln Asn Glu
210 215 220

Met Ile Glu Thr Glu Cys Phe Lys Glu Leu Asn Val Phe Gly Pro Asn
225 230 235 240

Gly Thr Leu Pro Pro Asp Leu Asn Arg Asn His Pro Pro Glu Pro Pro
245 250 255

Lys Lys Gly Leu Leu Gln Arg Leu Phe Lys Arg Gln His Gln Asn Asn
260 265 270

Ser Lys Ser Ser Pro Ser Ser Lys Thr Ser Phe Asn His His Ile Asn
275 280 285

Ser Asn His Val Ser Ser Asn Ser Thr Gly Ser Ser
290 295 300

<210> 117
<211> 169
<212> PRT
<213> Homo sapiens

<220>
<221> -
<222> (1) .. (169)
<223> "XAA" can be any amino acid

<400> 117

Met Arg Met Pro Arg Ala Arg Met Pro Xaa Pro Arg Ala His Ser Lys
1 5 10 15

Ala Gly Cys Pro Cys Gly Cys Pro Arg Asp Pro Leu Thr Leu Leu Ser
20 25 30

Pro Ser Gly His Ile Arg Ile Ser Asp Leu Gly Leu Ala Val Lys Ile
35 40 45

Pro Glu Gly Asp Leu Ile Arg Gly Arg Val Gly Thr Val Gly Tyr Met
50 55 60

Ala Pro Glu Val Leu Asn Asn Gln Arg Tyr Gly Leu Ser Pro Asp Tyr
65 70 75 80

Trp Gly Leu Gly Cys Leu Ile Tyr Glu Met Ile Glu Gly Gln Ser Pro
85 90 95

Phe Arg Gly Arg Lys Glu Lys Val Lys Arg Glu Glu Val Asp Arg Arg
100 105 110

Val Leu Glu Thr Glu Glu Val Tyr Ser His Lys Phe Ser Glu Glu Ala
115 120 125

Lys Ser Ile Cys Lys Met Val Ser Ser Trp Trp Pro Asp Ala Thr Leu
130 135 140

Lys Leu Val Ala Pro Ser Leu Gly Leu Ala Pro Val Cys Pro Gln Asn
145 150 155 160

Ser Lys Gln Ala Glu Gly Thr Gly Val
165

<210> 118

<211> 319

<212> PRT

<213> Homo sapiens

<400> 118

Met Ala Pro Phe Leu Arg Ile Ala Phe Asn Ser Tyr Glu Leu Gly Ser
1 5 10 15

Leu Gln Ala Glu Asp Glu Ala Asn Gln Pro Phe Cys Ala Val Lys Met
20 25 30

Lys Glu Ala Leu Ser Thr Glu Arg Gly Lys Thr Leu Val Gln Lys Lys
35 40 45

Pro Thr Met Tyr Pro Glu Trp Lys Ser Thr Phe Asp Ala His Ile Tyr
50 55 60

Glu Gly Arg Val Ile Gln Ile Val Leu Met Arg Ala Ala Glu Glu Pro
65 70 75 80

Val Ser Glu Val Thr Val Gly Val Ser Val Leu Ala Glu Arg Cys Lys
85 90 95

Lys Asn Asn Gly Lys Ala Glu Phe Trp Leu Asp Leu Gln Pro Gln Ala
100 105 110

Lys Val Leu Met Ser Val Gln Tyr Phe Leu Glu Asp Val Asp Cys Lys
115 120 125

Gln Ser Met Arg Ser Glu Asp Glu Ala Lys Phe Pro Thr Met Asn Arg
130 135 140

Arg Gly Ala Ile Lys Gln Ala Lys Ile His Tyr Ile Lys Asn His Glu
145 150 155 160

Phe Ile Ala Thr Phe Phe Gly Gln Pro Thr Phe Cys Ser Val Cys Lys
165 170 175

Asp Phe Val Trp Gly Leu Asn Lys Gln Gly Tyr Lys Cys Arg Gln Cys
180 185 190

Asn Ala Ala Ile His Lys Lys Cys Ile Asp Lys Ile Ile Gly Arg Cys
195 200 205

Thr Gly Thr Ala Ala Asn Ser Arg Asp Thr Ile Phe Gln Lys Glu Arg
210 215 220

Phe Asn Ile Asp Met Pro His Arg Phe Lys Val His Asn Tyr Met Ser
225 230 235 240

Pro Thr Phe Cys Asp His Cys Gly Ser Leu Leu Leu Pro Ala Pro His
245 250 255

Asp Lys His Gln Trp Asp Cys Gly Asp Phe Cys Cys Trp Pro Arg Pro
260 265 270

Cys Pro Gln Ser Val Leu Gly Cys Arg Leu Ala Gly Leu Ser Trp Tyr
275 280 285

Phe Leu Cys Glu Leu Cys Val Asn Leu Leu Phe Leu Cys Leu Arg Arg
290 295 300

Glu Ile Val Asn Pro Val Phe His Tyr Leu Asn Val Val Ile Tyr
305 310 315

<210> 119

<211> 236

<212> PRT

<213> Homo sapiens

<400> 119

Met Asp Glu Thr His Pro Gly Tyr Gly Lys Glu Val Asp Leu Glu Phe
1 5 10 15

Leu Val Ser Pro Ser Leu Pro Cys Leu Leu Ser Phe Ala Gly Ser Ala
20 25 30

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<210> 61
<211> 1662
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gacctagagg cacacgtccg gcagttgcag gagcggatgg agttgtgcgca ggcagaggga	300

Arg His Leu Val Pro Pro Asp Ser Asn Leu Phe Ser Lys Leu Trp Ala
35 40 45

Cys Gly Val Ile Leu Phe Thr Leu Leu Ala Gly Ser Pro Pro Phe Trp
50 55 60

His Arg Arg Gln Ile Leu Met Leu Arg Met Ile Met Glu Gly Gln Tyr
65 70 75 80

Gln Phe Ser Ser Pro Glu Trp Asp Asp Arg Ser Ser Thr Val Lys Asp
85 90 95

Leu Ile Ser Arg Leu Leu Gln Val Asp Pro Glu Ala Arg Leu Thr Ala
100 105 110

Glu Gln Ala Leu Gln His Pro Phe Phe Glu Arg Cys Glu Gly Ser Gln
115 120 125

Pro Trp Asn Leu Thr Pro Arg Gln Arg Phe Arg Val Ala Val Trp Thr
130 135 140

Val Leu Ala Ala Gly Arg Val Ala Leu Ser Thr His Arg Val Arg Pro
145 150 155 160

Leu Thr Lys Asn Ala Leu Leu Arg Asp Pro Tyr Ala Leu Arg Ser Val
165 170 175

Arg His Leu Ile Asp Asn Cys Ala Phe Arg Leu Tyr Gly His Trp Val
180 185 190

Lys Lys Gly Glu Gln Gln Asn Arg Ala Ala Leu Phe Gln His Arg Pro
195 200 205

Pro Gly Pro Phe Pro Ile Met Gly Pro Glu Glu Gly Asp Ser Ala
210 215 220

Ala Ile Thr Glu Asp Glu Ala Val Leu Val Leu Gly
225 230 235

<210> 120
<211> 572
<212> PRT
<213> Homo sapiens

<400> 120

Met Ala Phe Cys Ala Lys Met Arg Ser Ser Lys Lys Thr Glu Val Asn
1 5 10 15

Leu Glu Ala Pro Glu Pro Gly Val Glu Val Ile Phe Tyr Leu Ser Asp
20 25 30

Arg Glu Pro Leu Arg Leu Gly Ser Gly Glu Tyr Thr Ala Glu Glu Leu
35 40 45

Cys Ile Arg Ala Ala Gln Ala Cys Arg Ile Ser Pro Leu Cys His Asn

50

55

60

Leu Phe Ala Leu Tyr Asp Glu Asn Thr Lys Leu Trp Tyr Ala Pro Asn
65 70 75 80

Arg Thr Ile Thr Val Asp Asp Lys Met Ser Leu Arg Leu His Tyr Arg
85 90 95

Met Arg Phe Tyr Phe Thr Asn Trp His Gly Thr Asn Asp Asn Glu Gln
100 105 110

Ser Val Trp Arg His Ser Pro Lys Lys Gln Lys Asn Gly Tyr Glu Lys
115 120 125

Lys Lys Ile Pro Asp Ala Thr Pro Leu Leu Asp Ala Ser Ser Leu Glu
130 135 140

Tyr Leu Phe Ala Gln Gly Gln Tyr Asp Leu Val Lys Cys Leu Ala Pro
145 150 155 160

Ile Arg Asp Pro Lys Thr Glu Gln Asp Gly His Asp Ile Glu Asn Glu
165 170 175

Cys Leu Gly Met Ala Val Leu Ala Ile Ser His Tyr Ala Met Met Lys
180 185 190

Lys Met Gln Leu Pro Glu Leu Pro Lys Asp Ile Ser Tyr Lys Arg Tyr
195 200 205

Ile Pro Glu Thr Leu Asn Lys Ser Ile Arg Gln Arg Asn Leu Leu Thr
210 215 220

Arg Met Arg Ile Asn Asn Val Phe Lys Asp Phe Leu Lys Glu Phe Asn
225 230 235 240

Asn Lys Thr Ile Cys Asp Ser Ser Val Ser Thr His Asp Leu Lys Val
245 250 255

Lys Tyr Leu Ala Thr Leu Glu Thr Leu Thr Lys His Tyr Gly Ala Glu
260 265 270

Ile Phe Glu Thr Ser Met Leu Leu Ile Ser Ser Glu Asn Glu Met Asn
275 280 285

Trp Phe His Ser Asn Asp Gly Gly Asn Val Leu Tyr Tyr Glu Val Met
290 295 300

Val Thr Gly Asn Leu Gly Ile Gln Trp Arg His Lys Pro Asn Val Val
305 310 315 320

Ser Val Glu Lys Glu Lys Asn Lys Leu Lys Arg Lys Lys Leu Glu Asn
325 330 335

Lys Asp Lys Lys Asp Glu Glu Lys Asn Lys Ile Arg Glu Glu Trp Asn
340 345 350

Asn Phe Ser Phe Phe Pro Glu Ile Thr His Ile Val Ile Lys Glu Ser

355

360

365

Val Val Ser Ile Asn Lys Gln Asp Asn Lys Lys Met Glu Leu Lys Leu
370 375 380

Ser Ser His Glu Glu Ala Leu Ser Phe Val Ser Leu Val Asp Gly Tyr
385 390 395 400

Phe Arg Leu Thr Ala Asp Ala His His Tyr Leu Cys Thr Asp Val Ala
405 410 415

Pro Pro Leu Ile Val His Asn Ile Gln Asn Gly Cys His Gly Pro Ile
420 425 430

Cys Thr Glu Tyr Ala Ile Asn Lys Leu Arg Gln Glu Gly Ser Glu Glu
435 440 445

Gly Met Tyr Val Leu Arg Trp Ser Cys Thr Asp Phe Asp Asn Ile Leu
450 455 460

Met Thr Val Thr Cys Phe Glu Lys Ser Glu Gln Val Gln Gly Ala Gln
465 470 475 480

Lys Gln Phe Lys Asn Phe Gln Ile Glu Val Gln Lys Gly Arg Tyr Ser
485 490 495

Leu His Gly Ser Asp Arg Ser Phe Pro Ser Leu Gly Asp Leu Met Ser
500 505 510

His Leu Lys Lys Gln Ile Leu Arg Thr Asp Asn Ile Ser Phe Met Leu
515 520 525

Lys Arg Cys Cys Gln Pro Lys Pro Arg Gly Ser Leu Pro Val Pro Glu
530 535 540

Pro Gly Cys Ile Pro Ser Val Ile Ala Glu Thr Gln Ile Asp Gln Asn
545 550 555 560

Thr Leu Thr Asp Leu Asn Lys Val Asp Pro Pro Pro
565 570

<210> 121

<211> 311

<212> PRT

<213> Homo sapiens

<400> 121

Met Gly Cys Val Gln Cys Lys Asp Lys Glu Ala Thr Lys Leu Thr Glu
1 5 10 15

Glu Arg Asp Gly Ser Leu Asn Gln Ser Ser Gly Tyr Arg Tyr Gly Thr
20 25 30

Asp Pro Thr Pro Gln His Tyr Pro Ser Phe Gly Val Thr Ser Ile Pro
35 40 45

Asn Tyr Asn Asn Phe His Ala Ala Gly Gly Gln Gly Leu Thr Val Phe
50 55 60

Gly Gly Val Asn Ser Ser Ser His Thr Gly Thr Leu Arg Thr Arg Gly
65 70 75 80

Gly Thr Gly Val Thr Leu Phe Val Ala Leu Tyr Asp Tyr Glu Ala Arg
85 90 95

Thr Glu Asp Asp Leu Ser Phe His Lys Gly Glu Lys Phe Gln Ile Leu
100 105 110

Asn Ser Ser Glu Gly Asp Trp Trp Glu Ala Arg Ser Leu Thr Thr Gly
115 120 125

Glu Thr Gly Tyr Ile Pro Ser Asn Tyr Val Ala Pro Val Asp Ser Ile
130 135 140

Gln Ala Glu Glu Trp Tyr Phe Gly Lys Leu Gly Arg Lys Asp Ala Glu
145 150 155 160

Arg Gln Leu Leu Ser Phe Gly Asn Pro Arg Gly Thr Phe Leu Ile Arg
165 170 175

Glu Ser Glu Thr Thr Lys Gly Ala Tyr Ser Leu Ser Ile Arg Asp Trp
180 185 190

Asp Asp Met Lys Gly Asp His Val Lys His Tyr Lys Ile Arg Lys Leu
195 200 205

Asp Asn Gly Gly Tyr Tyr Ile Thr Thr Arg Ala Gln Phe Glu Thr Leu
210 215 220

Gln Gln Leu Val Gln His Tyr Ser Glu Arg Ala Ala Gly Leu Cys Cys
225 230 235 240

Arg Leu Val Val Pro Cys His Lys Gly Met Pro Arg Leu Thr Asp Leu
245 250 255

Ser Val Lys Thr Lys Asp Val Trp Glu Ile Pro Arg Glu Ser Leu Gln
260 265 270

Leu Ile Lys Arg Leu Gly Asn Gly Gln Phe Gly Glu Val Trp Met Gly
275 280 285

Met Leu Arg Leu Asn Tyr Ser Leu Ile Ser Phe Pro Val Trp Lys Ile
290 295 300

Pro Asn Thr Lys Asp Gly Arg
305 310

<210> 122
<211> 387
<212> PRT
<213> Homo sapiens

<400> 122

Met Gly Cys Val Gln Cys Lys Asp Lys Glu Ala Thr Lys Leu Thr Glu
1 5 10 15

Glu Arg Asp Gly Ser Leu Asn Gln Ser Ser Gly Tyr Arg Tyr Gly Thr
20 25 30

Asp Pro Thr Pro Gln His Tyr Pro Ser Phe Gly Val Thr Ser Ile Pro
35 40 45

Asn Tyr Asn Asn Phe His Ala Ala Gly Gly Gln Gly Leu Thr Val Phe
50 55 60

Gly Gly Val Asn Ser Ser His Thr Gly Thr Leu Arg Thr Arg Gly
65 70 75 80

Gly Thr Gly Val Thr Leu Phe Val Ala Leu Tyr Asp Tyr Glu Ala Arg
85 90 95

Thr Glu Asp Asp Leu Ser Phe His Lys Gly Glu Lys Phe Gln Ile Leu
100 105 110

Asn Ser Ser Glu Gly Asp Trp Trp Glu Ala Arg Ser Leu Thr Thr Gly
115 120 125

Glu Thr Gly Tyr Ile Pro Ser Asn Tyr Val Ala Pro Val Asp Ser Ile
130 135 140

Gln Ala Glu Glu Trp Tyr Phe Gly Lys Leu Gly Arg Lys Asp Ala Glu
145 150 155 160

Arg Gln Leu Leu Ser Phe Gly Asn Pro Arg Gly Thr Phe Leu Ile Arg
165 170 175

Glu Ser Glu Thr Thr Lys Gly Ala Tyr Ser Leu Ser Ile Arg Asp Trp
180 185 190

Asp Asp Met Lys Gly Asp His Val Lys His Tyr Lys Ile Arg Lys Leu
195 200 205

Asp Asn Gly Gly Tyr Tyr Ile Thr Thr Arg Ala Gln Phe Glu Thr Leu
210 215 220

Gln Gln Leu Val Gln His Tyr Ser Glu Arg Ala Ala Gly Leu Cys Cys
225 230 235 240

Arg Leu Val Val Pro Cys His Lys Gly Met Pro Arg Leu Thr Asp Leu
245 250 255

Ser Val Lys Thr Lys Asp Val Trp Glu Ile Pro Arg Glu Ser Leu Gln
260 265 270

Leu Ile Lys Arg Leu Gly Asn Gly Gln Phe Gly Glu Val Trp Met Gly
275 280 285

Thr Trp Asn Gly Asn Thr Lys Val Ala Ile Lys Thr Leu Lys Pro Gly
290 295 300

Thr Met Ser Pro Glu Ser Phe Leu Glu Glu Ala Gln Ile Met Lys Lys
305 310 315 320

Leu Lys His Asp Lys Leu Val Gln Leu Tyr Ala Val Val Ser Glu Glu
325 330 335

Pro Ile Tyr Ile Val Thr Glu Tyr Met Asn Lys Gly Trp Ala Thr Pro
340 345 350

Leu Leu Ser Pro Ala His Ser Ala Leu Arg Gly Cys Leu Gly Glu Arg
355 360 365

Asn Gly Ser Phe Leu Leu Ala Thr Phe Leu Val Ser Ala Trp Val Lys
370 375 380

Tyr Ser His
385

<210> 123
<211> 516
<212> PRT
<213> Homo sapiens

<400> 123

Met Arg Leu Glu Leu Pro Ala Gly His Trp Glu Arg Pro Asp Leu Glu
1 5 10 15

Leu Leu Glu Lys Ser Thr Gln Gln Gly Arg Ala Trp Asp Leu Glu Leu
20 25 30

Leu Glu Lys Gly Ala Gly Ser Leu Pro Leu Tyr Val Trp Lys Val Ser
35 40 45

Leu Ser Leu Leu Glu Leu His Lys Arg Arg Lys Ala Leu Thr Glu Pro
50 55 60

Glu Ala Arg Tyr Tyr Leu Arg Gln Ile Val Leu Gly Cys Gln Tyr Leu
65 70 75 80

His Arg Asn Arg Val Ile His Arg Asp Leu Lys Leu Gly Asn Leu Phe
85 90 95

Leu Asn Glu Asp Leu Glu Val Lys Ile Gly Asp Phe Gly Leu Ala Thr
100 105 110

Lys Val Glu Tyr Asp Gly Glu Arg Lys Lys Thr Leu Cys Gly Thr Pro
115 120 125

Asn Tyr Ile Ala Pro Glu Val Leu Ser Lys Lys Gly His Ser Phe Glu
130 135 140

Val Asp Val Trp Ser Ile Gly Cys Ile Met Tyr Thr Leu Leu Val Gly
145 150 155 160

Lys Pro Pro Phe Glu Thr Ser Cys Leu Lys Glu Thr Tyr Leu Arg Ile

165

170

175

Lys Lys Asn Glu Tyr Ser Ile Pro Lys His Ile Asn Pro Val Ala Ala
 180 185 190

Ser Leu Ile Gln Lys Met Leu Gln Thr Asp Pro Thr Ala Arg Pro Thr
 195 200 205

Ile Asn Glu Leu Leu Asn Asp Glu Phe Phe Thr Ser Gly Tyr Ile Pro
 210 215 220

Ala Arg Leu Pro Ile Thr Cys Leu Thr Ile Pro Pro Arg Phe Ser Ile
 225 230 235 240

Ala Pro Ser Ser Leu Asp Pro Ser Asn Arg Lys Pro Leu Thr Val Leu
 245 250 255

Asn Lys Gly Leu Glu Asn Pro Leu Pro Glu Arg Pro Arg Glu Lys Glu
 260 265 270

Glu Pro Val Val Arg Glu Thr Gly Glu Val Val Asp Cys His Leu Ser
 275 280 285

Asp Met Leu Gln Gln Leu His Ser Val Asn Ala Ser Lys Pro Ser Glu
 290 295 300

Arg Gly Leu Val Arg Gln Glu Glu Ala Glu Asp Pro Ala Cys Ile Pro
 305 310 315 320

Ile Phe Trp Val Ser Lys Trp Val Asp Tyr Ser Asp Lys Tyr Gly Leu
 325 330 335

Gly Tyr Gln Leu Cys Asp Asn Ser Val Gly Val Leu Phe Asn Asp Ser
 340 345 350

Thr Arg Leu Ile Leu Tyr Asn Asp Gly Asp Ser Leu Gln Tyr Ile Glu
 355 360 365

Arg Asp Gly Thr Glu Ser Tyr Leu Thr Val Ser Ser His Pro Asn Ser
 370 375 380

Leu Met Lys Lys Ile Thr Leu Leu Lys Tyr Phe Arg Asn Tyr Met Ser
 385 390 395 400

Glu His Leu Leu Lys Ala Gly Ala Asn Ile Thr Pro Arg Glu Gly Asp
 405 410 415

Glu Leu Ala Arg Leu Pro Tyr Leu Arg Thr Trp Phe Arg Thr Arg Ser
 420 425 430

Ala Ile Ile Leu His Leu Ser Asn Gly Ser Val Gln Ile Asn Phe Phe
 435 440 445

Gln Asp His Thr Lys Leu Ile Leu Cys Pro Leu Met Ala Ala Val Thr
 450 455 460

Tyr Ile Asp Glu Lys Arg Asp Phe Arg Thr Tyr Arg Leu Ser Leu Leu

465	470	475	480
Glu Glu Tyr Gly Cys Cys Lys Glu Leu Ala Ser Arg Leu Arg Tyr Ala			
485	490	495	
Arg Thr Met Val Asp Lys Leu Leu Ser Ser Arg Ser Ala Ser Asn Arg			
500	505	510	
Leu Lys Ala Ser			
515			
<210> 124			
<211> 171			
<212> PRT			
<213> Homo sapiens			
<220>			
<221> -			
<222> (1)..(171)			
<223> "XAA" can be any amino acid			
<400> 124			
Met Ala Leu Leu Pro Pro Phe Leu Ala Ser His Arg Leu Glu Val Ser			
1	5	10	15
Arg Asp Ser Gly Trp Leu Gly Gln Cys Trp Leu Gln Gly Val Trp Glu			
20	25	30	
Arg Xaa Pro His Ser Gly Leu Leu Tyr Pro Leu Gln His Pro Pro Ala			
35	40	45	
Glu Phe Ser Thr Tyr Leu Asn Phe Cys Arg Ser Leu Arg Phe Asp Asp			
50	55	60	
Lys Pro Asp Tyr Ser Tyr Leu Arg Gln Leu Phe Arg Asn Leu Phe His			
65	70	75	80
Arg Gln Gly Phe Ser Tyr Asp Tyr Val Phe Asp Trp Asn Met Leu Lys			
85	90	95	
Phe Gly Ala Ser Ser Ser Gln Ala Gln Pro Arg Asp Ser Pro Met Thr			
100	105	110	
Ala Lys Gly Pro Phe Cys Pro Arg Pro Cys Pro Cys Ala Gly Pro Thr			
115	120	125	
Tyr Ser Pro Thr Tyr Trp Cys Pro Ala Pro Leu Gly Thr Gln Ser Pro			
130	135	140	
Pro Asp Arg Pro Val Glu Glu Val Glu Glu Leu Ser Pro Gln Asn Tyr			
145	150	155	160
Trp Pro Val Val Trp Thr Pro Gly Pro His Phe			
165	170		

<210> 125
<211> 134
<212> PRT
<213> Homo sapiens

<400> 125

Met Ala Leu Leu Pro Pro Phe Leu Ala Ser His Arg Leu Glu Val Ser
1 5 10 15

Arg Asp Ser Gly Trp Leu Gly Gln Cys Trp Leu Gln Gly Val Trp Glu
20 25 30

Arg Gly Leu Thr Val Ala Phe Ser Ile Leu Cys Asn Thr Leu Gln Pro
35 40 45

Glu Phe Ser Thr Tyr Leu Asn Phe Cys Arg Ser Leu Arg Phe Asp Asp
50 55 60

Lys Pro Asp Tyr Ser Tyr Leu Arg Gln Leu Phe Arg Asn Leu Phe His
65 70 75 80

Arg Gln Gly Phe Ser Tyr Asp Tyr Val Phe Asp Trp Asn Met Leu Lys
85 90 95

Phe Gly Ala Ser Ser Ser Gln Ala Gln Pro Arg Asp Ser Pro Met Thr
100 105 110

Ala Lys Gly Pro Phe Cys Pro Arg Pro Cys Pro Cys Ala Gly Pro Thr
115 120 125

Tyr Ser Pro Thr Tyr Trp
130

<210> 126
<211> 233
<212> PRT
<213> Homo sapiens

<400> 126

Met Ala Leu Leu Pro Pro Phe Leu Ala Ser His Arg Leu Glu Val Ser
1 5 10 15

Arg Asp Ser Gly Trp Leu Gly Gln Cys Trp Leu Gln Gly Val Trp Glu
20 25 30

Arg Gly Leu Thr Val Ala Phe Ser Ile Leu Cys Asn Thr Leu Gln Pro
35 40 45

Glu Phe Ser Thr Tyr Leu Asn Phe Cys Arg Ser Leu Arg Phe Asp Asp
50 55 60

Lys Pro Asp Tyr Ser Tyr Leu Arg Gln Leu Phe Arg Asn Leu Phe His
65 70 75 80

Arg Gln Gly Phe Ser Tyr Asp Tyr Val Phe Asp Trp Asn Met Leu Lys

85

90

95

Phe Gly Gly Pro Leu Ser Cys Gln Pro Pro Ala Leu Pro Cys Gly Arg
100 105 110

Pro Gln Asp Glu Leu Gly Cys Ser Pro Glu Ser Arg Gly Cys Gly Pro
115 120 125

Gly Ala Ala Arg Thr Arg Thr Arg Gly Glu Asp Gly Ala Ala Thr Gly
130 135 140

Val Arg Asp Pro Ser Pro Ala Pro Trp Pro Thr His Gly Gly His Cys
145 150 155 160

Gln Pro Ala Pro Gln Cys Arg Arg Ala Arg Gly Phe His Ala Ser Leu
165 170 175

Pro His Pro Ala Gly Trp Gln Tyr Phe Ser Gln Ser Asp Leu Ala Gly
180 185 190

Arg Pro Gly Glu Glu Gly Glu Tyr Glu Ala Ala Gln Gly Cys Ala Arg
195 200 205

Gln Arg Leu Leu Leu Arg Pro His Trp Ala Ala Arg Gly Leu Pro Asp
210 215 220

Pro Ser Leu Thr Asp Lys Cys Ala Ile
225 230

<210> 127

<211> 243

<212> PRT

<213> Homo sapiens

<400> 127

Met Ala Ala Glu Leu Asn Lys Asn Lys Lys Ala Arg Ala Ala Glu Ala
1 5 10 15

Ala Arg Ala Ala Glu Ala Ala Lys Ala Ala Glu Ala Thr Lys Ala Ala
20 25 30

Glu Ala Ala Ala Lys Ala Ala Lys Ala Ser Asn Thr Ser Thr Pro Thr
35 40 45

Lys Gly Asn Thr Glu Thr Ser Ala Ser Ala Ser Gln Thr Asn His Val
50 55 60

Lys Asp Val Lys Lys Ile Lys Ile Glu His Ala Pro Ser Pro Ser Ser
65 70 75 80

Gly Gly Thr Leu Lys Asn Asp Lys Ala Lys Thr Lys Pro Pro Leu Gln
85 90 95

Val Thr Lys Val Glu Asn Asn Leu Ile Val Asp Lys Ala Thr Lys Lys
100 105 110

Ala Val Ile Val Gly Lys Glu Ser Lys Ser Ala Ala Thr Lys Glu Glu
115 120 125

Ser Val Ser Leu Lys Glu Lys Thr Lys Pro Leu Thr Pro Ser Ile Gly
130 135 140

Ala Lys Glu Lys Glu Gln His Val Ala Leu Val Thr Ser Thr Leu Pro
145 150 155 160

Pro Leu Pro Leu Pro Pro Met Leu Pro Glu Asp Lys Glu Ala Asp Ser
165 170 175

Leu Arg Gly Asn Ile Ser Val Lys Ala Val Lys Lys Glu Val Glu Lys
180 185 190

Lys Leu Arg Cys Leu Leu Ala Asp Leu Pro Leu Pro Pro Glu Leu Pro
195 200 205

Gly Gly Asp Asp Leu Ser Lys Ser Pro Glu Glu Lys Lys Thr Ala Thr
210 215 220

Gln Leu His Ser Lys Arg Arg Pro Lys Tyr Val Leu Ala Phe Tyr Leu
225 230 235 240

Leu Leu Asn

<210> 128

<211> 330

<212> PRT

<213> Homo sapiens

<400> 128

Met Ser Ala Lys Val Arg Leu Lys Lys Leu Glu Gln Leu Leu Asp
1 5 10 15

Gly Pro Trp Arg Asn Glu Ser Ala Leu Ser Val Glu Thr Leu Leu Asp
20 25 30

Val Leu Val Cys Leu Tyr Thr Glu Cys Ser His Ser Ala Leu Arg Arg
35 40 45

Asp Lys Tyr Val Ala Glu Phe Leu Glu Trp Ala Lys Pro Phe Thr Gln
50 55 60

Leu Val Lys Glu Met Gln Leu His Arg Glu Asp Phe Glu Ile Ile Lys
65 70 75 80

Val Ile Gly Arg Gly Ala Phe Gly Glu Val Ala Val Val Lys Met Lys
85 90 95

Asn Thr Glu Arg Ile Tyr Ala Met Lys Ile Leu Asn Lys Trp Glu Met
100 105 110

Leu Lys Arg Ala Glu Thr Ala Cys Phe Arg Glu Glu Arg Asp Val Leu
115 120 125

Val Asn Gly Asp Cys Gln Trp Ile Thr Ala Leu His Tyr Ala Phe Gln
130 135 140

Asp Glu Asn His Leu Tyr Leu Val Met Asp Tyr Tyr Val Gly Gly Asp
145 150 155 160

Leu Leu Thr Leu Leu Ser Lys Phe Glu Asp Lys Leu Pro Glu Asp Met
165 170 175

Ala Arg Phe Tyr Ile Gly Glu Met Val Leu Ala Ile Asp Ser Ile His
180 185 190

Gln Leu His Tyr Val His Arg Asp Ile Lys Pro Asp Asn Val Leu Leu
195 200 205

Asp Val Asn Gly His Ile Arg Leu Ala Asp Phe Gly Ser Cys Leu Lys
210 215 220

Met Asn Asp Asp Gly Thr Val Gln Ser Ser Val Ala Val Gly Thr Pro
225 230 235 240

Asp Tyr Ile Ser Pro Glu Ile Leu Gln Ala Met Glu Asp Gly Met Gly
245 250 255

Lys Tyr Gly Pro Glu Cys Asp Trp Trp Ser Leu Gly Val Cys Met Tyr
260 265 270

Glu Met Leu Tyr Gly Glu Thr Pro Phe Tyr Ala Glu Ser Leu Val Glu
275 280 285

Thr Tyr Gly Lys Ile Met Asn His Glu Glu Arg Phe Gln Phe Pro Ser
290 295 300

His Val Thr Asp Val Ser Glu Glu Ala Lys Asp Leu Ile Gln Arg Leu
305 310 315 320

Ser Cys Ile Gln Arg Thr Pro Tyr Leu Gln
325 330

<210> 129

<211> 246

<212> PRT

<213> Homo sapiens

<400> 129

Met Ser Ala Lys Val Arg Leu Lys Lys Leu Glu Gln Leu Leu Asp
1 5 10 15

Gly Pro Trp Arg Asn Glu Ser Ala Leu Ser Val Glu Thr Leu Leu Asp
20 25 30

Val Leu Val Cys Leu Tyr Thr Glu Cys Ser His Ser Ala Leu Arg Arg
35 40 45

Asp Lys Tyr Val Ala Glu Phe Leu Glu Trp Ala Lys Pro Phe Thr Gln

50

55

60

Leu Val Lys Glu Met Gln Leu His Arg Glu Asp Phe Glu Ile Ile Lys
65 70 75 80

Val Ile Gly Arg Gly Ala Phe Gly Glu Val Ala Val Val Lys Met Lys
85 90 95

Asn Thr Glu Arg Ile Tyr Ala Met Lys Ile Leu Asn Lys Trp Glu Met
100 105 110

Leu Lys Arg Ala Glu Thr Ala Cys Phe Arg Glu Glu Arg Asp Val Leu
115 120 125

Val Asn Gly Asp Cys Gln Trp Ile Thr Ala Leu His Tyr Ala Phe Gln
130 135 140

Asp Glu Asn His Leu Tyr Leu Val Met Asp Tyr Tyr Val Gly Gly Asp
145 150 155 160

Leu Leu Thr Leu Leu Ser Lys Phe Glu Asp Lys Leu Pro Glu Asp Met
165 170 175

Ala Arg Phe Tyr Ile Gly Glu Met Val Leu Ala Ile Asp Ser Ile His
180 185 190

Gln Leu His Tyr Val His Arg Asp Ile Lys Pro Asp Asn Val Leu Leu
195 200 205

Asp Val Asn Gly His Ile Arg Leu Ala Asp Phe Gly Ser Cys Leu Lys
210 215 220

Met Asn Asp Asp Gly Thr Val Gly Ile Phe Val Gly Asp Phe Pro Phe
225 230 235 240

Gly Phe Gly Phe Gly Ile
245

<210> 130

<211> 378

<212> PRT

<213> Homo sapiens

<400> 130

Met Glu Leu Arg Val Gly Asn Arg Tyr Arg Leu Gly Arg Lys Ile Gly
1 5 10 15

Ser Gly Ser Phe Gly Asp Ile Tyr Leu Val Gly Ile Pro Thr Ile Arg
20 25 30

Trp Cys Gly Ala Glu Gly Asp Tyr Asn Val Met Val Met Glu Leu Leu
35 40 45

Gly Pro Ser Leu Glu Asp Leu Phe Asn Phe Cys Ser Arg Lys Phe Ser
50 55 60

Leu Lys Thr Val Leu Leu Leu Ala Asp Gln Met Ile Ser Arg Ile Glu
65 70 75 80

Tyr Ile His Ser Lys Asn Phe Ile His Arg Asp Val Lys Pro Asp Asn
85 90 95

Phe Leu Met Gly Leu Gly Lys Gly Asn Leu Val Tyr Ile Ile Asp
100 105 110

Phe Gly Leu Ala Lys Lys Tyr Arg Asp Ala Arg Thr His Gln His Ile
115 120 125

Pro Tyr Arg Glu Asn Lys Asn Leu Thr Gly Thr Ala Arg Tyr Ala Ser
130 135 140

Ile Asn Thr His Leu Gly Ile Glu Gln Ser Arg Arg Asp Asp Leu Glu
145 150 155 160

Ser Leu Gly Tyr Val Leu Met Tyr Phe Asn Leu Gly Ser Leu Pro Trp
165 170 175

Gln Gly Leu Lys Ala Ala Thr Lys Arg Gln Lys Tyr Glu Arg Ile Ser
180 185 190

Glu Lys Lys Met Ser Thr Pro Ile Glu Val Leu Cys Lys Gly Tyr Pro
195 200 205

Ser Glu Phe Ala Thr Tyr Leu Asn Phe Cys Arg Ser Leu Arg Phe Asp
210 215 220

Asp Lys Pro Asp Tyr Ser Tyr Leu Arg Gln Leu Phe Arg Asn Leu Phe
225 230 235 240

His Arg Gln Gly Phe Ser Tyr Asp Tyr Val Phe Asp Trp Asn Met Leu
245 250 255

Lys Phe Gly Ala Ser Arg Ala Ala Asp Asp Ala Glu Arg Asp Ala Gly
260 265 270

Asp Arg Glu Glu Arg Leu Arg His Ser Arg Asn Pro Ala Thr Arg Gly
275 280 285

Leu Pro Ser Thr Ala Ser Gly Arg Leu Arg Gly Arg Arg Lys Val Ala
290 295 300

Pro Pro Thr Pro Leu Thr Pro Thr Ser His Thr Ala Asn Thr Ser Pro
305 310 315 320

Arg Pro Val Ser Gly Met Glu Arg Glu Arg Lys Val Ser Met Arg Leu
325 330 335

His Arg Gly Ala Pro Val Asn Ile Ser Ser Ser Asp Leu Thr Gly Arg
340 345 350

Gln Asp Thr Ser Arg Met Ser Thr Ser Gln Ile Pro Gly Arg Val Ala
355 360 365

Ser Ser Gly Leu Gln Ser Val Val His Arg
370 375

<210> 131
<211> 561
<212> PRT
<213> Homo sapiens

<400> 131

Met Val Glu Trp Trp Ser Ala Leu Thr Cys Pro Leu Gln Thr Phe Ala
1 5 10 15

Ala Pro Ser Phe Asp Asp Lys Ile Leu Glu Val Val Ala Val Phe Gly
20 25 30

Ser Met Gln Met Ala Val Ser Arg Val Ile Arg Leu Gln His His Arg
35 40 45

Ile Ala Gln Cys Arg Thr Val Lys Ile Ser Ile Leu Gly Asp Glu Gly
50 55 60

Val Pro Val Gln Val Asp Gly Glu Ala Trp Val Gln Pro Pro Gly Tyr
65 70 75 80

Ile Arg Ile Val His Lys Asn Arg Ala Gln Thr Leu Thr Arg Asp Arg
85 90 95

Ala Phe Glu Ser Thr Leu Lys Ser Trp Glu Asp Lys Gln Lys Cys Glu
100 105 110

Leu Pro Arg Pro Pro Ser Cys Ser Leu His Pro Glu Met Leu Ser Glu
115 120 125

Glu Glu Ala Thr Gln Met Asp Gln Phe Gly Gln Ala Ala Gly Val Leu
130 135 140

Ile His Ser Ile Arg Glu Ile Ala Gln Ser His Arg Asp Met Glu Gln
145 150 155 160

Glu Leu Ala His Ala Val Asn Ala Ser Ser Lys Ser Met Asp Arg Val
165 170 175

Tyr Gly Lys Pro Arg Thr Thr Glu Gly Leu Asn Cys Ser Phe Val Leu
180 185 190

Glu Met Val Asn Asn Phe Arg Ala Leu Arg Ser Glu Thr Glu Leu Leu
195 200 205

Leu Ser Gly Lys Met Ala Leu Gln Leu Asp Pro Pro Gln Lys Glu Gln
210 215 220

Leu Gly Ser Ala Leu Ala Glu Met Asp Arg Gln Leu Arg Arg Leu Ala
225 230 235 240

Asp Thr Pro Trp Leu Cys Gln Ser Ala Glu Pro Gly Asp Glu Glu Ser
245 250 255

Val Met Leu Asp Leu Ala Lys Arg Ser Arg Ser Gly Lys Phe Arg Leu
260 265 270

Val Thr Lys Phe Lys Lys Glu Lys Asn Asn Lys Asn Lys Glu Ala His
275 280 285

Ser Ser Leu Gly Ala Pro Val His Leu Trp Gly Thr Glu Glu Val Ala
290 295 300

Ala Trp Leu Glu His Leu Ser Leu Cys Glu Tyr Lys Asp Ile Phe Thr
305 310 315 320

Arg His Asp Ile Arg Gly Ser Glu Leu Leu His Leu Glu Arg Arg Asp
325 330 335

Leu Lys Asp Leu Gly Val Thr Lys Val Gly His Met Lys Arg Ile Leu
340 345 350

Cys Gly Ile Lys Glu Leu Ser Arg Ser Ala Pro Ala Val Glu Ala Gln
355 360 365

Pro Leu Ser Ser Gln Pro Val Ala Ser Thr Ser Pro Pro Pro Arg Pro
370 375 380

Ser Leu Arg Pro Leu Ser Leu Trp Pro Leu Arg Leu Leu Pro Leu Arg
385 390 395 400

Pro Trp Ala Asp Ala Ala Ala Arg Pro Leu Leu Met Val Leu Leu Pro
405 410 415

Leu Ser Ala Thr Glu Ser Leu Arg Asp Thr Val His Gln Ser Ser Gly
420 425 430

Val Ser Asn Ile Thr Thr Gln Leu Pro Leu Lys Gln His Phe Leu Gln
435 440 445

Leu Arg Val Thr Trp Gly Thr Cys Val Thr Ala Thr Gln Leu Ser Pro
450 455 460

Ala Cys Ala Val Gly Gln Gly Ile Gln Arg Arg Leu Ala Ser Trp Ala
465 470 475 480

Leu Leu Ala Trp Pro Arg Ala Trp Ile Val Pro Gly Ala Pro Leu Arg
485 490 495

Val Ser Phe Cys Gly Arg Thr Val Trp Leu Arg Leu Leu Ala Pro Ser
500 505 510

Gln Phe Ser Glu Thr Trp Leu Gly Pro Ser Thr Ala Ala Cys Lys Gly
515 520 525

Pro Cys Leu Leu Met Gln Leu Leu Asn Lys Asn Arg Ala Leu Ser
530 535 540

Trp Phe Glu Ser Ser Met Asp Val Ser Ser Leu Val Asp Cys Asn Leu
545 550 555 560

Thr

<210> 132
<211> 213
<212> PRT
<213> Homo sapiens

<400> 132

Met Ser Asp Val Ala Ile Val Lys Glu Gly Trp Leu His Lys Arg Gly
1 5 10 15

Glu Tyr Ile Lys Thr Trp Arg Pro Arg Tyr Phe Leu Leu Lys Asn Asp
20 25 30

Gly Thr Phe Ile Gly Tyr Lys Glu Arg Pro Gln Asp Val Asp Gln Arg
35 40 45

Glu Ala Pro Leu Asn Asn Phe Ser Val Ala Gln Cys Gln Leu Met Lys
50 55 60

Thr Glu Arg Pro Arg Pro Asn Thr Phe Ile Ile Arg Cys Leu Gln Trp
65 70 75 80

Thr Thr Val Ile Glu Arg Thr Phe His Val Glu Thr Pro Glu Glu Arg
85 90 95

Glu Glu Trp Thr Thr Ala Ile Gln Thr Val Ala Asp Gly Leu Lys Lys
100 105 110

Gln Glu Glu Glu Glu Met Asp Phe Arg Ser Gly Ser Pro Ser Asp Asn
115 120 125

Ser Gly Ala Glu Glu Met Glu Val Ser Leu Ala Lys Pro Lys His Arg
130 135 140

Val Ala Leu Gly Gly Arg Ala Gly Pro Ala His Val Ser Pro His Ser
145 150 155 160

Val Ser Gln Pro Pro Trp Ala Val Cys His Gln Leu Ser Val Ile Ser
165 170 175

Leu Gly Pro Trp Ala Ser Val Gln Pro Gly Gly Thr Arg Cys Asn Leu
180 185 190

Thr Met Val Cys Trp Pro Ala Pro Ser Pro Gly Gly Arg His Thr
195 200 205

Ala Ala Pro Gln His
210

<210> 133
<211> 425
<212> PRT
<213> Homo sapiens

<400> 133

Met Ile Val His Asp Asp Val Glu Ser Glu Pro Ala Met Thr Pro Ser
1 5 10 15

Lys Glu Gly Thr Leu Ile Val Arg Gln Thr Gln Ser Ala Ser Ser Thr
20 25 30

Leu Gln Lys His Lys Ser Ser Ser Phe Thr Pro Phe Ile Asp Pro
35 40 45

Arg Leu Leu Gln Ile Ser Pro Ser Ser Gly Thr Thr Val Thr Ser Val
50 55 60

Val Gly Phe Ser Cys Asp Gly Met Arg Pro Glu Ala Ile Arg Gln Asp
65 70 75 80

Pro Thr Arg Lys Gly Ser Val Val Asn Val Asn Pro Thr Asn Thr Arg
85 90 95

Pro Gln Ser Asp Thr Pro Glu Ile Arg Lys Tyr Lys Lys Arg Phe Asn
100 105 110

Ser Glu Ile Leu Cys Ala Ala Leu Trp Gly Val Asn Leu Leu Val Gly
115 120 125

Thr Glu Ser Gly Leu Met Leu Leu Asp Arg Ser Gly Gln Gly Lys Val
130 135 140

Tyr Pro Leu Ile Asn Arg Arg Arg Phe Gln Gln Met Asp Val Leu Glu
145 150 155 160

Gly Leu Asn Val Leu Val Thr Ile Ser Gly Lys Lys Asp Lys Leu Arg
165 170 175

Val Tyr Tyr Leu Ser Trp Leu Arg Asn Lys Ile Leu His Asn Asp Pro
180 185 190

Glu Val Glu Lys Lys Gln Gly Trp Thr Thr Val Gly Asp Leu Glu Gly
195 200 205

Cys Val His Tyr Lys Val Val Lys Tyr Glu Arg Ile Lys Phe Leu Val
210 215 220

Ile Ala Leu Lys Ser Ser Val Glu Val Tyr Ala Trp Ala Pro Lys Pro
225 230 235 240

Tyr His Lys Phe Met Ala Phe Lys Ser Phe Gly Glu Leu Val His Lys
245 250 255

Pro Leu Leu Val Asp Leu Thr Val Glu Glu Gly Gln Arg Leu Lys Val
260 265 270

Ile Tyr Gly Ser Cys Ala Gly Phe His Ala Val Asp Val Asp Ser Gly
275 280 285

Ser Val Tyr Asp Ile Tyr Leu Pro Thr His Ile Gln Cys Ser Ile Lys
290 295 300

Pro His Ala Ile Ile Ile Leu Pro Asn Thr Asp Gly Met Glu Leu Leu
305 310 315 320

Val Cys Tyr Glu Asp Glu Gly Val Tyr Val Asn Thr Tyr Gly Arg Ile
325 330 335

Thr Lys Asp Val Val Leu Gln Trp Gly Glu Met Pro Thr Ser Val Ala
340 345 350

Tyr Ile Arg Ser Asn Gln Thr Met Gly Trp Gly Glu Lys Ala Ile Glu
355 360 365

Ile Arg Ser Val Glu Thr Gly His Leu Asp Gly Val Phe Met His Lys
370 375 380

Arg Ala Gln Arg Leu Lys Phe Leu Cys Glu Arg Asn Asp Lys Val Phe
385 390 395 400

Phe Ala Ser Val Arg Ser Gly Gly Ser Ser Gln Val Tyr Phe Met Thr
405 410 415

Leu Gly Arg Thr Ser Leu Leu Ser Trp
420 425

<210> 134
<211> 515
<212> PRT
<213> Homo sapiens

<400> 134

Met Ala Ser Arg Thr Pro Arg Asn Cys Ala Val Leu Lys Gly Glu Val
1 5 10 15

Asp Leu Thr Ala Leu Ala Lys Glu Leu Arg Ala Val Glu Asp Val Arg
20 25 30

Pro Pro His Lys Val Thr Asp Tyr Ser Ser Ser Ser Glu Glu Ser Gly
35 40 45

Thr Thr Asp Glu Glu Asp Asp Val Glu Gln Glu Gly Ala Asp Glu
50 55 60

Ser Thr Ser Gly Pro Glu Asp Thr Arg Ala Ala Ser Ser Leu Asn Leu
65 70 75 80

Ser Asn Gly Glu Thr Glu Ser Val Lys Thr Met Ile Val His Asp Asp
85 90 95

Val Glu Ser Glu Pro Ala Met Thr Pro Ser Lys Glu Gly Thr Leu Ile
100 105 110

Val Arg Gln Thr Gln Ser Ala Ser Ser Thr Leu Gln Lys His Lys Ser
115 120 125

Ser Ser Ser Phe Thr Pro Phe Ile Asp Pro Arg Leu Leu Gln Ile Ser
130 135 140

Pro Ser Ser Gly Thr Thr Val Thr Ser Val Val Gly Phe Ser Cys Asp
145 150 155 160

Gly Met Arg Pro Glu Ala Ile Arg Gln Asp Pro Thr Arg Lys Gly Ser
165 170 175

Val Val Asn Val Asn Pro Thr Asn Thr Arg Pro Gln Ser Asp Thr Pro
180 185 190

Glu Ile Arg Lys Tyr Lys Lys Arg Phe Asn Ser Glu Ile Leu Cys Ala
195 200 205

Ala Leu Trp Gly Val Asn Leu Leu Val Gly Thr Glu Ser Gly Leu Met
210 215 220

Leu Leu Asp Arg Ser Gly Gln Gly Lys Val Tyr Pro Leu Ile Asn Arg
225 230 235 240

Arg Arg Phe Gln Gln Met Asp Val Leu Glu Gly Leu Asn Val Leu Val
245 250 255

Thr Ile Ser Gly Lys Lys Asp Lys Leu Arg Val Tyr Tyr Leu Ser Trp
260 265 270

Leu Arg Asn Lys Ile Leu His Asn Asp Pro Glu Val Glu Lys Lys Gln
275 280 285

Gly Trp Thr Thr Val Gly Asp Leu Glu Gly Cys Val His Tyr Lys Val
290 295 300

Val Lys Tyr Glu Arg Ile Lys Phe Leu Val Ile Ala Leu Lys Ser Ser
305 310 315 320

Val Glu Val Tyr Ala Trp Ala Pro Lys Pro Tyr His Lys Phe Met Ala
325 330 335

Phe Lys Ser Phe Gly Glu Leu Val His Lys Pro Leu Leu Val Asp Leu
340 345 350

Thr Val Glu Glu Gly Gln Arg Leu Lys Val Ile Tyr Gly Ser Cys Ala
355 360 365

Gly Phe His Ala Val Asp Val Asp Ser Gly Ser Val Tyr Asp Ile Tyr
370 375 380

Leu Pro Thr His Ile Gln Cys Ser Ile Lys Pro His Ala Ile Ile Ile
385 390 395 400

Leu Pro Asn Thr Asp Gly Met Glu Leu Leu Val Cys Tyr Glu Asp Glu
405 410 415

Gly Val Tyr Val Asn Thr Tyr Gly Arg Ile Thr Lys Asp Val Val Leu
420 425 430

Gln Trp Gly Glu Met Pro Thr Ser Val Ala Tyr Ile Arg Ser Asn Gln
435 440 445

Thr Met Gly Trp Gly Glu Lys Ala Ile Glu Ile Arg Ser Val Glu Thr
450 455 460

Gly His Leu Asp Gly Val Phe Met His Lys Arg Ala Gln Arg Leu Lys
465 470 475 480

Phe Leu Cys Glu Arg Asn Asp Lys Val Phe Phe Ala Ser Val Arg Ser
485 490 495

Gly Gly Ser Ser Gln Val Tyr Phe Met Thr Leu Gly Arg Thr Ser Leu
500 505 510

Leu Ser Trp
515

<210> 135
<211> 468
<212> PRT
<213> Homo sapiens

<400> 135

Met Ser Ala Arg Val Gln Leu Thr Lys Ser Val Pro Ala Ile Met Arg
1 5 10 15

Ala Met Ala Leu Arg Phe Ala Phe Thr Ser Cys Gln Ile Ser Tyr Ser
20 25 30

Lys Ala Ile Pro Pro Pro Leu Pro Pro Pro Pro Pro His Pro Pro
35 40 45

Ala Ser Arg His Pro Pro Cys Pro His Arg His Pro Arg Asp Lys Leu
50 55 60

Thr Ala Asn Glu Thr Gln Ser Ala Ser Ser Thr Leu Gln Lys His Lys
65 70 75 80

Ser Ser Ser Ser Phe Thr Pro Phe Ile Asp Pro Arg Leu Leu Gln Ile
85 90 95

Ser Pro Ser Ser Gly Thr Thr Val Thr Ser Val Val Gly Phe Ser Cys
100 105 110

Asp Gly Met Arg Pro Glu Ala Ile Arg Gln Asp Pro Thr Arg Lys Gly
115 120 125

Ser Val Val Asn Val Asn Pro Thr Asn Thr Arg Pro Gln Ser Asp Thr
130 135 140

Pro Glu Ile Arg Lys Tyr Lys Arg Phe Asn Ser Glu Ile Leu Cys
145 150 155 160

Ala Ala Leu Trp Gly Val Asn Leu Leu Val Gly Thr Glu Ser Gly Leu

165

170

175

Met Leu Leu Asp Arg Ser Gly Gln Gly Lys Val Tyr Pro Leu Ile Asn
180 185 190

Arg Arg Arg Phe Gln Gln Met Asp Val Leu Glu Gly Leu Asn Val Leu
195 200 205

Val Thr Ile Ser Gly Lys Lys Asp Lys Leu Arg Val Tyr Tyr Leu Ser
210 215 220

Trp Leu Arg Asn Lys Ile Leu His Asn Asp Pro Glu Val Glu Lys Lys
225 230 235 240

Gln Gly Trp Thr Thr Val Gly Asp Leu Glu Gly Cys Val His Tyr Lys
245 250 255

Val Val Lys Tyr Glu Arg Ile Lys Phe Leu Val Ile Ala Leu Lys Ser
260 265 270

Ser Val Glu Val Tyr Ala Trp Ala Pro Lys Pro Tyr His Lys Phe Met
275 280 285

Ala Phe Lys Ser Phe Gly Glu Leu Val His Lys Pro Leu Leu Val Asp
290 295 300

Leu Thr Val Glu Glu Gly Gln Arg Leu Lys Val Ile Tyr Gly Ser Cys
305 310 315 320

Ala Gly Phe His Ala Val Asp Val Asp Ser Gly Ser Val Tyr Asp Ile
325 330 335

Tyr Leu Pro Thr His Ile Gln Cys Ser Ile Lys Pro His Ala Ile Ile
340 345 350

Ile Leu Pro Asn Thr Asp Gly Met Glu Leu Leu Val Cys Tyr Glu Asp
355 360 365

Glu Gly Val Tyr Val Asn Thr Tyr Gly Arg Ile Thr Lys Asp Val Val
370 375 380

Leu Gln Trp Gly Glu Met Pro Thr Ser Val Ala Tyr Ile Arg Ser Asn
385 390 395 400

Gln Thr Met Gly Trp Gly Glu Lys Ala Ile Glu Ile Arg Ser Val Glu
405 410 415

Thr Gly His Leu Asp Gly Val Phe Met His Lys Arg Ala Gln Arg Leu
420 425 430

Lys Phe Leu Cys Glu Arg Asn Asp Lys Val Phe Phe Ala Ser Val Arg
435 440 445

Ser Gly Gly Ser Ser Gln Val Tyr Phe Met Thr Leu Gly Arg Thr Ser
450 455 460

Leu Leu Ser Trp

465

<210> 136
<211> 666
<212> PRT
<213> Homo sapiens

<220>
<221> -
<222> (1)..(666)
<223> "XAA" can be any amino acid

<400> 136

Met Asp Cys Gln Leu Ser Ile Leu Leu Leu Ser Cys Ser Val Leu
1 5 10 15

Asp Ser Phe Gly Glu Leu Ile Pro Gln Pro Ser Asn Glu Val Asn Leu
20 25 30

Leu Asp Ser Lys Thr Ile Gln Gly Glu Leu Gly Trp Ile Ser Tyr Pro
35 40 45

Ser His Gly Trp Glu Glu Ile Ser Gly Val Asp Glu His Tyr Thr Pro
50 55 60

Ile Arg Thr Tyr Gln Val Cys Asn Val Met Asp His Ser Gln Asn Asn
65 70 75 80

Trp Leu Arg Thr Asn Trp Val Pro Arg Asn Ser Ala Gln Lys Ile Tyr
85 90 95

Val Glu Leu Lys Phe Thr Leu Arg Asp Cys Asn Ser Ile Pro Leu Val
100 105 110

Leu Gly Thr Cys Lys Glu Thr Phe Asn Leu Tyr Tyr Met Glu Ser Asp
115 120 125

Asp Asp His Gly Val Lys Phe Arg Glu His Gln Phe Thr Lys Ile Asp
130 135 140

Thr Ile Ala Ala Asp Glu Ser Phe Thr Gln Met Asp Leu Gly Asp Arg
145 150 155 160

Ile Leu Lys Leu Asn Thr Glu Ile Arg Glu Val Gly Pro Val Asn Lys
165 170 175

Lys Gly Phe Tyr Leu Ala Phe Gln Asp Val Gly Ala Cys Val Ala Leu
180 185 190

Val Ser Val Arg Val Tyr Phe Lys Lys Cys Pro Phe Thr Val Lys Asn
195 200 205

Leu Ala Met Phe Pro Asp Thr Val Pro Met Asp Ser Gln Ser Leu Val
210 215 220

Glu Val Arg Gly Ser Cys Val Asn Asn Ser Lys Glu Glu Asp Pro Pro
225 230 235 240

Arg Met Tyr Cys Ser Thr Glu Gly Glu Trp Leu Val Pro Ile Gly Lys
245 250 255

Cys Ser Cys Asn Ala Gly Tyr Glu Glu Arg Gly Phe Met Cys Gln Ala
260 265 270

Cys Arg Pro Gly Phe Tyr Lys Ala Leu Asp Gly Asn Met Lys Cys Ala
275 280 285

Lys Cys Pro Pro His Ser Ser Thr Gln Glu Asp Gly Ser Met Asn Cys
290 295 300

Arg Cys Glu Asn Asn Tyr Phe Arg Ala Asp Lys Asp Pro Pro Ser Met
305 310 315 320

Ala Cys Thr Arg Pro Pro Ser Ser Pro Arg Asn Val Ile Ser Asn Ile
325 330 335

Asn Glu Thr Ser Val Ile Leu Asp Trp Ser Trp Pro Leu Asp Thr Gly
340 345 350

Gly Arg Lys Asp Val Thr Phe Asn Ile Ile Cys Lys Lys Cys Gly Trp
355 360 365

Asn Ile Lys Gln Cys Glu Pro Cys Ser Pro Asn Val Arg Phe Leu Pro
370 375 380

Arg Gln Phe Gly Leu Thr Asn Thr Thr Val Thr Val Thr Asp Leu Leu
385 390 395 400

Ala His Thr Asn Tyr Thr Phe Glu Ile Asp Ala Val Asn Gly Val Ser
405 410 415

Glu Leu Ser Ser Pro Pro Arg Gln Phe Ala Ala Val Ser Ile Thr Thr
420 425 430

Asn Gln Ala Ala Pro Ser Pro Val Leu Thr Ile Lys Lys Asp Arg Thr
435 440 445

Ser Arg Asn Ser Ile Ser Leu Ser Trp Gln Glu Pro Glu His Pro Asn
450 455 460

Gly Ile Ile Leu Asp Tyr Glu Val Lys Tyr Glu Lys Gln Glu Gln
465 470 475 480

Glu Thr Ser Tyr Thr Ile Leu Arg Ala Arg Gly Thr Asn Val Thr Ile
485 490 495

Ser Ser Leu Lys Pro Asp Thr Ile Tyr Val Phe Gln Ile Arg Ala Arg
500 505 510

Thr Ala Ala Gly Tyr Gly Thr Asn Ser Arg Lys Phe Glu Phe Glu Thr
515 520 525

Ser Pro Asp Ser Phe Ser Ile Ser Gly Glu Ser Ser Gln Val Val Met
530 535 540

Ile Ala Ile Ser Ala Ala Val Ala Ile Ile Leu Leu Thr Val Val Ile
545 550 555 560

Tyr Val Leu Ile Gly Arg Phe Cys Gly Tyr Lys Ser Lys His Gly Ala
565 570 575

Asp Glu Lys Arg Leu His Phe Gly Asn Gly His Leu Lys Leu Pro Gly
580 585 590

Leu Arg Thr Tyr Val Asp Pro His Thr Tyr Glu Asp Pro Thr Gln Ala
595 600 605

Val His Glu Phe Ala Lys Glu Leu Asp Ala Thr Asn Ile Ser Ile Asp
610 615 620

Lys Val Val Gly Ala Val Leu Thr Ser Glu Gln Leu His Asp Ala Glu
625 630 635 640

Xaa Phe Ser Leu Ala Gly Phe Asn Val Ser Ser Gln Gly Val His Phe
645 650 655

Ser Pro Ala Arg Ser Leu Pro Val Ala Asn
660 665

<210> 137
<211> 458
<212> PRT
<213> Homo sapiens

<400> 137

Met Lys Tyr Thr Phe Trp Gly Trp Val Ala Val Val Lys Leu Lys Asn
1 5 10 15

Ala Asp Lys Val Phe Ala Met Lys Ile Leu Asn Lys Trp Glu Met Leu
20 25 30

Lys Arg Ala Glu Thr Ala Cys Phe Arg Glu Glu Arg Asp Val Leu Val
35 40 45

Asn Gly Asp Asn Lys Trp Ile Thr Thr Leu His Tyr Ala Phe Gln Asp
50 55 60

Asp Asn Asn Leu Tyr Leu Val Met Asp Tyr Tyr Val Gly Gly Asp Leu
65 70 75 80

Leu Thr Leu Leu Ser Lys Phe Glu Asp Arg Leu Pro Glu Asp Met Ala
85 90 95

Arg Phe Tyr Leu Ala Glu Met Val Ile Ala Ile Asp Ser Val His Gln
100 105 110

Leu His Tyr Val His Arg Asp Ile Lys Pro Asp Asn Ile Leu Met Asp
115 120 125

Met Asn Gly His Ile Arg Leu Ala Asp Phe Gly Ser Cys Leu Lys Leu
130 135 140

Met Glu Asp Gly Thr Val Gln Ser Ser Val Ala Val Gly Thr Pro Asp
145 150 155 160

Tyr Ile Ser Pro Glu Ile Leu Gln Ala Met Glu Asp Gly Lys Gly Arg
165 170 175

Tyr Gly Pro Glu Cys Asp Trp Trp Ser Leu Gly Val Cys Met Tyr Glu
180 185 190

Met Leu Tyr Gly Glu Thr Pro Phe Tyr Ala Glu Ser Leu Val Glu Thr
195 200 205

Tyr Gly Lys Ile Met Asn His Lys Glu Arg Phe Gln Phe Pro Ala Gln
210 215 220

Val Thr Asp Val Ser Glu Asn Ala Lys Asp Leu Ile Arg Arg Leu Ile
225 230 235 240

Cys Ser Arg Glu His Arg Leu Gly Gln Asn Gly Ile Glu Asp Phe Lys
245 250 255

Lys His Pro Phe Phe Ser Gly Ile Asp Trp Asp Asn Ile Arg Asn Cys
260 265 270

Glu Ala Pro Tyr Ile Pro Glu Val Ser Ser Pro Thr Asp Thr Ser Asn
275 280 285

Phe Asp Val Asp Asp Asp Cys Leu Lys Asn Ser Glu Thr Met Pro Pro
290 295 300

Pro Thr His Thr Ala Phe Ser Gly His His Leu Pro Phe Val Gly Phe
305 310 315 320

Thr Tyr Thr Ser Ser Cys Val Leu Ser Asp Arg Ser Cys Leu Arg Val
325 330 335

Thr Ala Gly Pro Thr Ser Leu Asp Leu Asp Val Asn Val Gln Arg Thr
340 345 350

Leu Asp Asn Asn Leu Ala Thr Glu Ala Tyr Glu Arg Arg Ile Lys Arg
355 360 365

Leu Glu Gln Glu Lys Leu Glu Leu Ser Arg Lys Leu Gln Glu Ser Thr
370 375 380

Gln Thr Val Gln Ala Leu Gln Tyr Ser Thr Val Asp Gly Pro Leu Thr
385 390 395 400

Ala Ser Lys Asp Leu Glu Ile Lys Asn Leu Lys Glu Glu Ile Glu Lys
405 410 415

Leu Arg Lys Gln Val Thr Glu Ser Ser His Leu Glu Gln Gln Leu Glu
420 425 430

Glu Ala Asn Ala Val Arg Gln Glu Leu Asp Asp Ala Phe Arg Gln Ile
435 440 445

Lys Ala Tyr Glu Lys Gln Ile Lys Thr Leu
450 455

<210> 138
<211> 262
<212> PRT
<213> Homo sapiens

<400> 138

Met Glu Val Val Asp Pro Gln Gln Leu Gly Met Phe Thr Glu Gly Glu
1 5 10 15

Leu Met Ser Val Gly Met Asp Thr Phe Ile His Arg Ile Asp Ser Thr
20 25 30

Glu Val Ile Tyr Gln Pro Arg Arg Lys Arg Ala Lys Leu Ile Gly Lys
35 40 45

Tyr Leu Met Gly Asp Leu Leu Gly Glu Gly Ser Tyr Gly Lys Val Lys
50 55 60

Glu Val Leu Asp Ser Glu Thr Leu Cys Arg Arg Ala Val Lys Ile Leu
65 70 75 80

Lys Lys Lys Leu Arg Arg Ile Pro Asn Gly Glu Ala Asn Val Lys
85 90 95

Lys Glu Ile Gln Leu Leu Arg Arg Leu Arg His Lys Asn Val Ile Gln
100 105 110

Leu Val Asp Val Leu Tyr Asn Glu Glu Lys Gln Lys Met Tyr Met Val
115 120 125

Met Glu Tyr Cys Val Cys Gly Met Gln Glu Met Leu Asp Ser Val Pro
130 135 140

Glu Lys Arg Phe Pro Val Cys Gln Ala His Gly Ser Pro Ser Arg Arg
145 150 155 160

Gly Gly Arg His Ala Ser Val Pro Thr Thr Pro Gln Asp Leu Arg Ser
165 170 175

Ala Leu Gln Gly Arg Ala Gly Gly Gln Gln Gly Pro Gly Ala Ala Leu
180 185 190

Pro Pro Arg Pro Pro Gly Ser Ala Arg Gly Leu Leu Thr Ser Gln Pro
195 200 205

Arg Ala Glu Pro Ser Arg Ala Gly Val Gly Gly Arg Arg Pro Pro
210 215 220

Cys Thr Leu Cys Gly Asp Tyr Trp Pro Arg Pro Trp Pro Arg Ala Pro

225 230 235 240
Gln Gly Ala Gln Arg Arg Pro Ala Ala Pro Pro Gln Thr Ser Trp Arg
 245 250 255
Val Trp Arg Pro Gly Ser
 260

<210> 139
<211> 203
<212> PRT
<213> Homo sapiens

<400> 139

Met Glu Val Val Asp Pro Gln Gln Leu Gly Met Phe Thr Glu Gly Glu
1 5 10 15

Leu Met Ser Val Gly Met Asp Thr Phe Ile His Arg Ile Asp Ser Thr
20 25 30

Glu Val Ile Tyr Gln Pro Arg Arg Lys Arg Ala Lys Leu Ile Gly Lys
35 40 45

Tyr Leu Met Gly Asp Leu Leu Gly Glu Gly Ser Tyr Gly Lys Val Lys
50 55 60

Glu Val Leu Asp Ser Glu Thr Leu Cys Arg Arg Ala Val Lys Ile Leu
65 70 75 80

Lys Lys Lys Lys Leu Arg Arg Ile Pro Asn Gly Glu Ala Asn Val Lys
85 90 95

Lys Glu Ile Gln Leu Leu Arg Arg Leu Arg His Lys Asn Val Ile Gln
100 105 110

Leu Val Asp Val Leu Tyr Asn Glu Glu Lys Gln Lys Met Tyr Met Val
115 120 125

Met Glu Tyr Cys Val Cys Gly Met Gln Glu Met Leu Asp Ser Val Pro
130 135 140

Glu Lys Arg Phe Pro Val Cys Gln Ala His Gly Tyr Phe Cys Gln Leu
145 150 155 160

Ile Asp Gly Leu Glu Tyr Leu His Ser Gln Gly Ile Val His Lys Asp
165 170 175

Ile Lys Pro Gly Asn Leu Leu Thr Thr Gly Gly Thr Leu Lys Ile
180 185 190

Ser Asp Leu Gly Val Ala Glu Val Gly Thr Cys
195 200

<210> 140
<211> 244
<212> PRT

<213> Homo sapiens

<400> 140

Met Asp Arg Glu Thr Thr Pro Leu Gly Leu Leu Trp Leu Ile Gln Val
1 5 10 15

Ile Pro Ser Lys Leu Leu Pro Ser Leu Gln Val Lys Asp Phe Leu Ser
20 25 30

Gln Leu Arg Ser Ser Asn Arg Arg Phe Ser Ile Pro Glu Ser Gly Gln
35 40 45

Gly Gly Thr Glu Met Asp Gly Phe Arg Arg Thr Ile Glu Asn Gln His
50 55 60

Ser Arg Asn Asp Val Met Val Ser Glu Trp Leu Asn Lys Leu Asn Leu
65 70 75 80

Glu Glu Pro Pro Ser Ser Val Pro Lys Lys Cys Pro Ser Leu Thr Lys
85 90 95

Arg Ser Arg Ala Gln Glu Glu Gln Val Pro Gln Ala Trp Thr Ala Gly
100 105 110

Thr Ser Ser Asp Ser Met Ala Gln Pro Pro Gln Thr Pro Glu Thr Ser
115 120 125

Thr Phe Arg Asn Gln Met Pro Ser Pro Thr Ser Thr Gly Thr Pro Ser
130 135 140

Pro Gly Pro Arg Gly Asn Gln Gly Ala Glu Arg Gln Gly Met Asn Trp
145 150 155 160

Ser Cys Arg Thr Pro Glu Pro Asn Pro Val Thr Gly Arg Pro Leu Val
165 170 175

Asn Ile Tyr Asn Cys Ser Gly Val Gln Val Gly Asp Asn Asn Tyr Leu
180 185 190

Thr Met Gln Gln Thr Thr Ala Leu Pro Thr Trp Gly Leu Ala Pro Ser
195 200 205

Gly Lys Gly Arg Gly Leu Gln His Pro Pro Pro Val Gly Ser Gln Glu
210 215 220

Gly Pro Lys Asp Pro Glu Ala Trp Ser Arg Pro Gln Gly Trp Tyr Asn
225 230 235 240

His Ser Gly Lys

<210> 141

<211> 222

<212> PRT

<213> Homo sapiens

<400> 141

Met Val Lys Leu Tyr Leu Tyr Gln Lys Asn Val Lys Ile Ala Ile Phe
1 5 10 15

Asp Leu Lys Ser Arg Gln Asn Phe Phe Val Tyr Phe Arg Glu Glu Gln
20 25 30

Ala Arg Glu Leu Tyr Arg Arg Leu Arg Glu Lys Pro Arg Asp Gln Arg
35 40 45

Thr Glu Gly Asp Ser Gln Glu Met Val Arg Leu Leu Leu Gln Ala Ile
50 55 60

Gln Ser Phe Glu Lys Lys Val Arg Val Ile Tyr Thr Gln Leu Ser Lys
65 70 75 80

Thr Val Val Cys Lys Gln Lys Ala Leu Glu Leu Leu Pro Lys Val Glu
85 90 95

Glu Val Val Ser Leu Met Asn Glu Asp Glu Lys Thr Val Val Arg Leu
100 105 110

Gln Glu Lys Arg Gln Lys Glu Leu Trp Asn Leu Leu Lys Ile Ala Cys
115 120 125

Ser Lys Val Arg Gly Pro Val Ser Gly Ser Pro Asp Ser Met Asn Ala
130 135 140

Ser Arg Leu Ser Gln Pro Gly Gln Leu Met Ser Gln Pro Ser Thr Ala
145 150 155 160

Ser Asn Ser Leu Pro Glu Pro Ala Lys Lys Ser Glu Glu Leu Val Ala
165 170 175

Glu Ala His Asn Leu Cys Thr Leu Leu Glu Asn Ala Ile Gln Asp Thr
180 185 190

Val Arg Glu Gln Asp Gln Ser Phe Thr Ala Leu Asp Trp Ser Trp Leu
195 200 205

Gln Thr Glu Glu Glu His Ser Cys Leu Glu Gln Ala Ser
210 215 220

<210> 142

<211> 409

<212> PRT

<213> Homo sapiens

<400> 142

Met Arg Leu Thr Leu Leu Cys Cys Thr Trp Arg Glu Glu Arg Met Gly
1 5 10 15

Glu Glu Gly Ser Glu Leu Pro Val Cys Ala Ser Cys Gly Gln Arg Ile
20 25 30

Tyr Asp Gly Gln Tyr Leu Gln Ala Leu Asn Ala Asp Trp His Ala Asp
35 40 45

Cys Phe Arg Cys Cys Asp Cys Ser Ala Ser Leu Ser His Gln Tyr Tyr
50 55 60

Glu Lys Asp Gly Gln Leu Phe Cys Lys Lys Asp Tyr Trp Ala Arg Tyr
65 70 75 80

Gly Glu Ser Cys His Gly Cys Ser Glu Gln Ile Thr Lys Gly Leu Val
85 90 95

Met Val Ala Gly Glu Leu Lys Tyr His Pro Glu Cys Phe Ile Cys Leu
100 105 110

Thr Cys Gly Thr Phe Ile Gly Asp Gly Asp Thr Tyr Thr Leu Val Glu
115 120 125

His Ser Lys Leu Tyr Cys Gly His Cys Tyr Tyr Gln Thr Val Val Thr
130 135 140

Pro Val Ile Glu Gln Ile Leu Pro Asp Ser Pro Gly Ser His Leu Pro
145 150 155 160

His Thr Val Thr Leu Val Ser Ile Pro Ala Ser Ser His Gly Lys Arg
165 170 175

Gly Leu Ser Val Ser Ile Asp Pro Pro His Gly Pro Pro Gly Cys Gly
180 185 190

Thr Glu His Ser His Thr Val Arg Val Gln Gly Val Asp Pro Gly Cys
195 200 205

Met Ser Pro Asp Val Lys Asn Ser Ile His Val Gly Asp Arg Ile Leu
210 215 220

Glu Ile Asn Gly Thr Pro Ile Arg Asn Val Pro Leu Asp Glu Ile Asp
225 230 235 240

Leu Leu Ile Gln Glu Thr Ser Arg Leu Leu Gln Leu Thr Leu Glu His
245 250 255

Asp Pro His Asp Thr Leu Gly His Gly Leu Gly Pro Glu Thr Ser Pro
260 265 270

Leu Ser Ser Pro Ala Tyr Thr Pro Ser Gly Glu Ala Gly Ser Ser Ala
275 280 285

Arg Gln Lys Pro Val Leu Arg Ser Cys Ser Ile Asp Arg Ser Pro Gly
290 295 300

Ala Gly Ser Leu Gly Ser Pro Ala Ser Gln Arg Lys Asp Leu Gly Arg
305 310 315 320

Ser Glu Ser Leu Arg Val Val Cys Arg Pro His Arg Ile Phe Arg Pro
325 330 335

Ser Asp Leu Ile His Gly Glu Val Leu Gly Lys Gly Cys Phe Gly Gln
340 345 350

Ala Ile Lys Val Gln Ser Met Pro Gly Ser Gln Leu Asp Ser Leu Gly
355 360 365

Gly Thr Pro Pro Ser Ser Phe Leu Pro Ser Leu Trp Lys His Ser Gly
370 375 380

Arg Gly Ile Trp Leu Ser Asp Ser Leu Ala Ser Ala Leu Ser Ser Leu
385 390 395 400

Gly Leu Leu Glu Leu Ile Arg Asn Arg
405

<210> 143

<211> 305

<212> PRT

<213> Homo sapiens

<400> 143

Met Arg Leu Thr Leu Leu Cys Cys Thr Trp Arg Glu Glu Arg Met Gly
1 5 10 15

Glu Glu Gly Ser Glu Leu Pro Val Cys Ala Ser Cys Gly Gln Arg Ile
20 25 30

Tyr Asp Gly Gln Tyr Leu Gln Ala Leu Asn Ala Asp Trp His Ala Asp
35 40 45

Cys Phe Arg Cys Cys Asp Cys Ser Ala Ser Leu Ser His Gln Tyr Tyr
50 55 60

Glu Lys Asp Gly Gln Leu Phe Cys Lys Lys Asp Tyr Trp Ala Arg Tyr
65 70 75 80

Gly Glu Ser Cys His Gly Cys Ser Glu Gln Ile Thr Lys Gly Leu Val
85 90 95

Met Val Ala Gly Glu Leu Lys Tyr His Pro Glu Cys Phe Ile Cys Leu
100 105 110

Thr Cys Gly Thr Phe Ile Gly Asp Gly Asp Thr Tyr Thr Leu Val Glu
115 120 125

His Ser Lys Leu Tyr Cys Gly His Cys Tyr Tyr Gln Thr Val Val Thr
130 135 140

Pro Val Ile Glu Gln Ile Leu Pro Asp Ser Pro Gly Ser His Leu Pro
145 150 155 160

His Thr Val Thr Leu Val Ser Ile Pro Ala Ser Ser His Gly Lys Arg
165 170 175

Gly Leu Ser Val Ser Ile Asp Pro Pro His Gly Pro Pro Gly Cys Gly
180 185 190

Thr Glu His Ser His Thr Val Arg Val Gln Gly Val Asp Pro Gly Cys
195 200 205

Met Ser Pro Asp Val Lys Asn Ser Ile His Val Gly Asp Arg Ile Leu
210 215 220

Glu Ile Asn Gly Thr Pro Ile Arg Asn Val Pro Leu Asp Glu Ile Asp
225 230 235 240

Leu Leu Ile Gln Glu Thr Ser Arg Leu Leu Gln Leu Thr Leu Glu His
245 250 255

Asp Pro His Asp Thr Leu Gly His Gly Leu Gly Pro Glu Thr Ser Pro
260 265 270

Leu Ser Ser Pro Ala Tyr Thr Pro Ser Gly Glu Ala Gly Ser Ser Ala
275 280 285

Arg Gln Lys Pro Val Phe Ala Arg Thr Trp Val Ala Leu Ser Pro Ser
290 295 300

Ala
305

<210> 144
<211> 780
<212> PRT
<213> Homo sapiens

<400> 144

Met Ala Ser Asp Ala Val Gln Ser Glu Pro Arg Ser Trp Ser Leu Leu
1 5 10 15

Glu Gln Leu Gly Leu Ala Gly Ala Asp Leu Ala Ala Pro Gly Val Gln
20 25 30

Gln Gln Leu Glu Leu Glu Arg Glu Arg Leu Arg Arg Glu Ile Arg Lys
35 40 45

Glu Leu Lys Leu Lys Glu Gly Ala Glu Asn Leu Arg Arg Ala Thr Thr
50 55 60

Asp Leu Gly Arg Ser Leu Gly Pro Val Glu Leu Leu Leu Arg Gly Ser
65 70 75 80

Ser Arg Arg Leu Asp Leu Leu His Gln Gln Leu Gln Glu Leu His Ala
85 90 95

His Val Val Leu Pro Asp Pro Ala Ala Thr His Asp Gly Pro Gln Ser
100 105 110

Pro Gly Ala Gly Gly Pro Thr Cys Ser Ala Thr Asn Leu Ser Arg Val
115 120 125

Ala Gly Leu Glu Lys Gln Leu Ala Ile Glu Leu Lys Val Lys Gln Gly

130 135 140

Ala Glu Asn Met Ile Gln Thr Tyr Ser Asn Gly Ser Thr Lys Asp Arg
145 150 155 160

Lys Leu Leu Leu Thr Ala Gln Gln Met Leu Gln Asp Ser Lys Thr Lys
165 170 175

Ile Asp Ile Ile Arg Met Gln Leu Arg Arg Ala Leu Gln Ala Asp Gln
180 185 190

Leu Glu Asn Gln Ala Ala Pro Asp Asp Thr Gln Gly Ser Pro Asp Leu
195 200 205

Gly Ala Val Glu Leu Arg Ile Glu Glu Leu Arg His His Phe Arg Val
210 215 220

Glu His Ala Val Ala Glu Gly Ala Lys Asn Val Leu Arg Leu Leu Ser
225 230 235 240

Ala Ala Lys Ala Pro Asp Arg Lys Ala Val Ser Glu Ala Gln Glu Lys
245 250 255

Leu Thr Glu Ser Asn Gln Lys Leu Gly Leu Leu Arg Glu Ala Leu Glu
260 265 270

Arg Arg Leu Gly Glu Leu Pro Ala Asp His Pro Lys Gly Arg Leu Leu
275 280 285

Arg Glu Glu Leu Ala Ala Ser Ser Ala Ala Phe Ser Thr Arg Leu
290 295 300

Ala Gly Pro Phe Pro Ala Thr His Tyr Ser Thr Leu Cys Lys Pro Ala
305 310 315 320

Pro Leu Thr Gly Thr Leu Glu Val Arg Val Val Gly Cys Arg Asp Leu
325 330 335

Pro Glu Thr Ile Pro Trp Asn Pro Thr Pro Ser Met Gly Pro Gly
340 345 350

Thr Pro Asp Ser Arg Pro Pro Phe Leu Ser Arg Pro Ala Arg Gly Leu
355 360 365

Tyr Ser Arg Ser Gly Ser Leu Ser Gly Arg Ser Ser Leu Lys Ala Glu
370 375 380

Ala Glu Asn Thr Ser Glu Val Ser Thr Val Leu Lys Leu Asp Asn Thr
385 390 395 400

Val Val Gly Gln Thr Ser Trp Lys Pro Cys Gly Pro Asn Ala Trp Asp
405 410 415

Gln Ser Phe Thr Leu Glu Leu Glu Arg Ala Arg Glu Leu Glu Leu Ala
420 425 430

Val Phe Trp Arg Asp Gln Arg Gly Leu Cys Ala Leu Lys Phe Leu Lys

435

440

445

Leu Glu Asp Phe Leu Asp Asn Glu Arg His Glu Val Gln Leu Asp Met
450 455 460

Glu Pro Gln Gly Cys Leu Val Ala Glu Val Thr Phe Arg Asn Pro Val
465 470 475 480

Ile Glu Arg Ile Pro Arg Leu Arg Arg Gln Lys Lys Ile Phe Ser Lys
485 490 495

Gln Gln Gly Lys Ala Phe Gln Arg Ala Arg Gln Met Asn Ile Asp Val
500 505 510

Ala Thr Trp Val Arg Leu Leu Arg Arg Leu Ile Pro Asn Ala Thr Gly
515 520 525

Thr Gly Thr Phe Ser Pro Gly Ala Ser Pro Gly Ser Glu Ala Arg Thr
530 535 540

Thr Gly Asp Ile Ser Val Glu Lys Leu Asn Leu Gly Thr Asp Ser Asp
545 550 555 560

Ser Ser Pro Gln Lys Ser Ser Arg Asp Pro Pro Ser Ser Pro Ser Ser
565 570 575

Leu Ser Ser Pro Ile Gln Glu Ser Thr Ala Pro Glu Leu Pro Ser Glu
580 585 590

Thr Gln Glu Thr Pro Gly Pro Ala Leu Cys Ser Pro Leu Arg Lys Ser
595 600 605

Pro Leu Thr Leu Glu Asp Phe Lys Phe Leu Ala Val Leu Gly Arg Gly
610 615 620

His Phe Gly Lys Val Leu Leu Ser Glu Phe Arg Pro Ser Gly Glu Leu
625 630 635 640

Phe Ala Ile Lys Ala Leu Lys Lys Gly Asp Ile Val Ala Arg Asp Glu
645 650 655

Val Glu Ser Leu Met Cys Glu Lys Arg Ile Leu Ala Ala Val Thr Ser
660 665 670

Ala Gly His Pro Phe Leu Val Asn Leu Phe Gly Cys Phe Gln Thr Pro
675 680 685

Glu His Val Cys Phe Val Met Glu Tyr Ser Ala Gly Gly Asp Leu Met
690 695 700

Leu His Ile His Ser Asp Val Phe Ser Glu Pro Arg Ala Ile Phe Tyr
705 710 715 720

Ser Ala Cys Arg Leu Pro Pro Phe Val Pro Thr Leu Ser Gly Arg
725 730 735

Thr Asp Val Ser Asn Phe Asp Glu Glu Phe Thr Gly Glu Ala Pro Thr

740

745

750

Leu Ser Pro Pro Arg Asp Ala Arg Pro Leu Thr Ala Ala Glu Gln Ala
755 760 765

Ala Phe Leu Asp Phe Asp Phe Val Ala Gly Gly Cys
770 775 780

<210> 145

<211> 401

<212> PRT

<213> Homo sapiens

<400> 145

Met Ala Ser Asp Ala Val Gln Ser Glu Pro Arg Ser Trp Ser Leu Leu
1 5 10 15

Glu Gln Leu Gly Leu Ala Gly Ala Asp Leu Ala Ala Pro Gly Val Gln
20 25 30

Gln Gln Leu Glu Leu Glu Arg Glu Arg Leu Arg Arg Glu Ile Arg Lys
35 40 45

Glu Leu Lys Leu Lys Glu Gly Ala Glu Asn Leu Arg Arg Ala Thr Thr
50 55 60

Asp Leu Gly Arg Ser Leu Gly Pro Val Glu Leu Leu Leu Arg Gly Ser
65 70 75 80

Ser Arg Arg Leu Asp Leu Leu His Gln Gln Leu Gln Glu Leu His Ala
85 90 95

His Val Val Leu Pro Asp Pro Ala Ala Thr His Asp Gly Pro Gln Ser
100 105 110

Pro Gly Ala Gly Gly Pro Thr Cys Ser Ala Thr Asn Leu Ser Arg Val
115 120 125

Ala Gly Leu Glu Lys Gln Leu Ala Ile Glu Leu Lys Val Lys Gln Gly
130 135 140

Ala Glu Asn Met Ile Gln Thr Tyr Ser Asn Gly Ser Thr Lys Asp Arg
145 150 155 160

Lys Leu Leu Leu Thr Ala Gln Gln Met Leu Gln Asp Ser Lys Thr Lys
165 170 175

Ile Asp Ile Ile Arg Met Gln Leu Arg Arg Ala Leu Gln Ala Asp Gln
180 185 190

Leu Glu Asn Gln Ala Ala Pro Asp Asp Thr Gln Gly Ser Pro Asp Leu
195 200 205

Gly Ala Val Glu Leu Arg Ile Glu Glu Leu Arg His His Phe Arg Val
210 215 220

Glu His Ala Val Ala Glu Gly Ala Lys Asn Val Leu Arg Leu Leu Ser
225 230 235 240

Ala Ala Lys Ala Pro Asp Arg Lys Ala Val Ser Glu Ala Gln Glu Lys
245 250 255

Leu Thr Glu Ser Asn Gln Lys Leu Gly Leu Leu Arg Glu Ala Leu Glu
260 265 270

Arg Arg Leu Gly Glu Leu Pro Ala Asp His Pro Lys Gly Arg Leu Leu
275 280 285

Arg Glu Glu Leu Ala Ala Ser Ser Ala Ala Phe Ser Thr Arg Leu
290 295 300

Ala Gly Pro Phe Pro Ala Thr His Tyr Ser Thr Leu Cys Lys Pro Ala
305 310 315 320

Pro Leu Thr Gly Thr Leu Glu Val Arg Val Val Gly Cys Arg Asp Leu
325 330 335

Pro Glu Thr Ile Pro Trp Asn Pro Thr Pro Ser Met Gly Gly Pro Gly
340 345 350

Thr Pro Asp Ser Arg Pro Pro Phe Leu Ser Arg Pro Ala Arg Gly Leu
355 360 365

Tyr Ser Arg Ser Gly Ser Leu Ser Gly Arg Ser Ser Leu Lys Ala Glu
370 375 380

Ala Glu Asn Thr Ser Glu Val Ser Thr Val Leu Lys Leu Asp Asn Thr
385 390 395 400

His

<210> 146
<211> 96
<212> PRT
<213> Homo sapiens

<400> 146

Met Gln Ser Phe Leu Val Glu Gly Arg Phe Lys His Glu Met Phe Glu
1 5 10 15

Lys Val Phe Ala Glu Glu Arg Asn Gly Gly Gln Arg Leu Leu Cys Ala
20 25 30

Thr Asp Val Pro Ile Arg Thr Val Ser Ser Ala Ala Ser Gln Gly Leu
35 40 45

His Met Gln Asn Asp Asp Ala Cys Leu Gly Ala Ala Ser Pro Ser Ala
50 55 60

Ala Ser Trp Ser Arg Arg Ser Ala Glu Ser Lys Val Ser Leu Cys Trp
65 70 75 80

Lys Leu Lys Trp Lys Glu Asp Leu Val Trp Phe Tyr Ser Gln Ser His
85 90 95

<210> 147

<211> 333

<212> PRT

<213> Homo sapiens

<400> 147

Met His Arg Tyr Phe Glu Ser Pro Arg Arg Leu Leu Pro Val His Phe
1 5 10 15

Cys Cys Cys Gln Trp Arg Gly Gly Val Asp Phe Glu Cys Leu Leu
20 25 30

Gly Gly Val Trp Asp Arg Cys Arg Lys Val Leu Arg Ala Gln Glu Cys
35 40 45

Glu Trp Pro Arg His Leu Pro Ser Ala Cys Leu Leu Ser Ser Ala Cys
50 55 60

Arg Gly Gln Pro Glu Arg Arg Ala Ala Val Val Gly Ala Gln Asp Pro
65 70 75 80

Thr Glu Pro Pro Arg Leu Ser Arg Ser Leu Ser Gly Ala Ser Pro Phe
85 90 95

Leu Gly Glu Thr Lys Gln Glu Thr Leu Thr Asn Ile Ser Ala Val Asn
100 105 110

Tyr Asp Phe Asp Glu Glu Tyr Phe Ser Asn Thr Ser Glu Leu Ala Lys
115 120 125

Asp Phe Ile Arg Arg Leu Leu Val Lys Asp Pro Lys Arg Arg Met Thr
130 135 140

Ile Ala Gln Ser Leu Glu His Ser Trp Ile Lys Ala Ile Arg Arg Arg
145 150 155 160

Asn Val Arg Gly Glu Asp Ser Gly Arg Lys Pro Glu Arg Arg Arg Leu
165 170 175

Lys Thr Thr Arg Leu Lys Glu Tyr Thr Ile Lys Ser His Ser Ser Leu
180 185 190

Pro Pro Asn Asn Ser Tyr Ala Asp Phe Glu Arg Phe Ser Lys Val Leu
195 200 205

Glu Glu Ala Ala Ala Ala Glu Glu Gly Leu Arg Glu Leu Gln Arg Ser
210 215 220

Arg Arg Leu Cys His Glu Asp Val Glu Ala Leu Ala Ala Ile Tyr Glu
225 230 235 240

Glu Lys Glu Ala Trp Tyr Arg Glu Glu Ser Asp Ser Leu Gly Gln Asp

245

250

255

Leu Arg Arg Leu Arg Gln Glu Leu Leu Lys Thr Glu Ala Leu Lys Arg
260 265 270

Gln Ala Gln Glu Glu Ala Lys Gly Ala Leu Leu Gly Thr Ser Gly Leu
275 280 285

Lys Arg Arg Phe Ser Arg Leu Glu Asn Arg Tyr Glu Ala Leu Ala Lys
290 295 300

Gln Val Ala Ser Glu Met Arg Phe Val Gln Asp Leu Val Arg Ala Leu
305 310 315 320

Glu Gln Glu Lys Leu Gln Gly Val Glu Cys Gly Leu Arg
325 330

<210> 148

<211> 131

<212> PRT

<213> Homo sapiens

<400> 148

Met Leu Lys Glu Phe Leu Glu Ile Pro Phe Pro Thr Ser Pro Glu Cys
1 5 10 15

Thr Leu Gln Pro Lys Ser Gln Gln Pro Thr Gly Lys Glu Ala Glu Glu
20 25 30

His Pro Thr Ser Ala Pro Leu Thr His Ser Leu Leu Pro Pro Thr Pro
35 40 45

Leu Trp Val Val Ser His Phe Ile Phe Asp Phe Arg Gly Glu Thr Ala
50 55 60

Leu His Lys Ala Ala Cys Gln Arg Asn Arg Ala Val Cys Gln Leu Leu
65 70 75 80

Val Asp Ala Gly Ala Ser Leu Arg Lys Thr Asp Ser Lys Gly Lys Thr
85 90 95

Pro Gln Glu Arg Ala Gln Gln Ala Gly Asp Pro Asp Leu Ala Ala Tyr
100 105 110

Leu Glu Ser Arg Gln Asn Tyr Lys Val Ile Gly His Glu Asp Leu Glu
115 120 125

Thr Ala Val
130

<210> 149

<211> 272

<212> PRT

<213> Homo sapiens

<400> 149

Met Arg Gly Ala Ala Arg Leu Gly Arg Pro Gly Arg Ser Cys Leu Pro
1 5 10 15

Gly Pro Ala Leu Arg Ala Pro Pro Arg Pro Pro Leu Leu Leu Leu Leu
20 25 30

Ala Leu Leu Pro Leu Leu Pro Ala Pro Gly Ala Ala Ala Ala Pro Ala
35 40 45

Pro Arg Pro Pro Glu Leu Gln Ser Ala Ser Ala Gly Pro Ser Val Ser
50 55 60

Leu Tyr Leu Ser Glu Asp Glu Val Arg Arg Leu Ile Gly Leu Asp Ala
65 70 75 80

Glu Leu Tyr Tyr Val Arg Asn Asp Leu Ile Ser His Tyr Ala Leu Ser
85 90 95

Phe Ser Leu Leu Val Pro Ser Glu Thr Asn Phe Leu His Phe Thr Trp
100 105 110

His Ala Lys Ser Lys Val Glu Tyr Lys Leu Gly Phe Gln Val Asp Asn
115 120 125

Val Leu Ala Met Asp Met Pro Gln Val Asn Ile Ser Val Gln Gly Glu
130 135 140

Val Pro Arg Thr Leu Ser Val Phe Arg Val Glu Leu Ser Cys Thr Gly
145 150 155 160

Lys Val Asp Ser Glu Val Met Ile Leu Met Gln Leu Asn Leu Thr Val
165 170 175

Asn Ser Ser Lys Asn Phe Thr Val Leu Asn Phe Lys Arg Arg Lys Met
180 185 190

Cys Tyr Lys Lys Leu Glu Glu Val Lys Thr Ser Ala Leu Asp Lys Asn
195 200 205

Thr Ser Arg Thr Ile Tyr Asp Pro Val His Ala Ala Pro Thr Thr Ser
210 215 220

Thr Arg Val Phe Tyr Ile Ser Val Gly Val Cys Cys Ala Val Ile Phe
225 230 235 240

Leu Val Ala Ile Ile Leu Ala Val Leu His Leu His Ser Met Lys Arg
245 250 255

Ile Glu Leu Asp Asp Arg Tyr Cys Thr Tyr Phe Gly Lys Glu Lys Lys
260 265 270

<210> 150
<211> 344
<212> PRT
<213> Homo sapiens

<400> 150

Met Pro Gln Val Asn Ile Ser Val Gln Gly Glu Val Pro Arg Thr Leu
1 5 10 15

Ser Val Phe Arg Val Glu Leu Ser Cys Thr Gly Lys Val Asp Ser Glu
20 25 30

Val Met Ile Leu Met Gln Leu Asn Leu Thr Val Asn Ser Ser Lys Asn
35 40 45

Phe Thr Val Leu Asn Phe Lys Arg Arg Lys Met Cys Tyr Lys Lys Leu
50 55 60

Glu Glu Val Lys Thr Ser Ala Leu Asp Lys Asn Thr Ser Arg Thr Ile
65 70 75 80

Tyr Asp Pro Val His Ala Ala Pro Thr Thr Ser Thr Arg Val Phe Tyr
85 90 95

Ile Ser Val Gly Val Cys Cys Ala Val Ile Phe Leu Val Ala Ile Ile
100 105 110

Leu Ala Val Leu His Leu His Ser Met Lys Arg Ile Glu Leu Asp Asp
115 120 125

Ser Ile Ser Ala Ser Ser Ser Gln Gly Leu Ser Gln Pro Ser Thr
130 135 140

Gln Thr Thr Gln Tyr Leu Arg Ala Asp Thr Pro Asn Asn Ala Thr Pro
145 150 155 160

Ile Thr Ser Ser Tyr Tyr Pro Thr Leu Arg Ile Glu Lys Asn Asp Leu
165 170 175

Arg Ser Val Thr Leu Leu Glu Ala Lys Gly Lys Val Lys Asp Ile Ala
180 185 190

Ile Ser Arg Glu Arg Ile Thr Leu Lys Asp Val Leu Gln Glu Gly Thr
195 200 205

Phe Gly Arg Ile Phe His Gly Ile Leu Ile Asp Glu Lys Asp Pro Asn
210 215 220

Lys Glu Lys Gln Ala Phe Val Lys Thr Val Lys Asp Gln Ala Ser Glu
225 230 235 240

Ile Gln Val Thr Met Met Leu Thr Glu Ser Cys Lys Leu Arg Gly Leu
245 250 255

His His Arg Asn Leu Leu Pro Ile Thr His Val Cys Ile Glu Glu Gly
260 265 270

Glu Lys Pro Met Val Ile Leu Pro Tyr Met Asn Trp Gly Asn Leu Lys
275 280 285

Leu Phe Leu Arg Gln Cys Lys Leu Val Glu Ala Asn Asn Pro Gln Ala

290

295

300

Ile Ser Gln Gln Asp Leu Val His Met Ala Ile Gln Ile Ala Cys Gly
305 310 315 320

Met Ser Tyr Leu Ala Arg Arg Glu Val Ile His Lys Asp Leu Ala Ala
325 330 335

Arg Asn Cys Val Gly Pro Leu Glu
340

<210> 151

<211> 141

<212> PRT

<213> Homo sapiens

<400> 151

Met Glu Ala Ile Arg Thr Asp Asn Gln Asn Phe Ala Ser Gln Leu Arg
1 5 10 15

Glu Ala Glu Ala Arg Asn Arg Asp Leu Glu Ala His Val Arg Gln Leu
20 25 30

Gln Glu Arg Met Glu Leu Leu Gln Ala Glu Gly Ala Thr Ala Val Thr
35 40 45

Gly Val Pro Ser Pro Arg Ala Thr Asp Pro Pro Ser His Leu Asp Gly
50 55 60

Pro Pro Ala Val Ala Val Gly Gln Cys Pro Leu Val Gly Pro Gly Pro
65 70 75 80

Met His Arg Arg His Leu Leu Leu Pro Ala Arg Val Pro Arg Pro Gly
85 90 95

Leu Ser Glu Ala Leu Ser Leu Leu Leu Phe Ala Val Val Leu Ser Arg
100 105 110

Ala Ala Ala Leu Gly Cys Ile Gly Leu Val Ala His Ala Gly Gln Leu
115 120 125

Thr Ala Val Trp Arg Arg Pro Gly Ala Ala Arg Ala Pro
130 135 140

<210> 152

<211> 106

<212> PRT

<213> Homo sapiens

<400> 152

Met Glu Leu Leu Gln Ala Glu Gly Ala Thr Ala Val Thr Gly Val Pro
1 5 10 15

Ser Pro Arg Ala Thr Asp Pro Pro Ser His Leu Asp Gly Pro Pro Ala
20 25 30

Val Ala Val Gly Gln Cys Pro Leu Val Gly Pro Gly Pro Met His Arg
35 40 45

Arg His Leu Leu Leu Pro Ala Arg Val Pro Arg Pro Gly Leu Ser Glu
50 55 60

Ala Leu Ser Leu Leu Leu Phe Ala Val Val Leu Ser Arg Ala Ala Ala
65 70 75 80

Leu Gly Cys Ile Gly Leu Val Ala His Ala Gly Gln Leu Thr Ala Val
85 90 95

Trp Arg Arg Pro Gly Ala Ala Arg Ala Pro
100 105

<210> 153
<211> 50
<212> PRT
<213> Homo sapiens

<400> 153

Met Val Asn Leu Ser His Glu Asp Phe Glu Phe Ile Ser Gly Thr Arg
1 5 10 15

Met Arg Lys Leu Ala Arg Glu Gly Gln Lys Pro Pro Glu Gly Phe Met
20 25 30

Ala Pro Lys Ala Trp Thr Val Leu Thr Glu Tyr Tyr Lys Ser Leu Glu
35 40 45

Lys Ala
50

<210> 154
<211> 238
<212> PRT
<213> Homo sapiens

<400> 154

Met Ala Arg Thr Thr Ser Gln Leu Tyr Asp Ala Val Pro Ile Gln Ser
1 5 10 15

Ser Val Val Leu Cys Ser Cys Pro Ser Pro Ser Met Val Arg Thr Gln
20 25 30

Thr Glu Ser Ser Thr Pro Pro Gly Ile Pro Gly Gly Ser Arg Gln Gly
35 40 45

Pro Ala Met Asp Gly Thr Ala Ala Glu Pro Arg Pro Gly Ala Gly Ser
50 55 60

Leu Gln His Ala Gln Pro Pro Pro Gln Pro Arg Lys Lys Arg Pro Glu
65 70 75 80

Asp Phe Lys Phe Gly Lys Ile Leu Gly Glu Gly Ser Phe Ser Thr Val
85 90 95

Val Leu Ala Arg Glu Leu Ala Thr Ser Arg Glu Tyr Ala Ile Lys Ile
100 105 110

Leu Glu Lys Arg His Ile Ile Lys Glu Asn Lys Val Pro Tyr Val Thr
115 120 125

Arg Glu Arg Asp Val Met Ser Arg Leu Asp His Pro Phe Phe Val Lys
130 135 140

Leu Tyr Phe Thr Phe Gln Asp Asp Glu Lys Leu Tyr Phe Gly Leu Ser
145 150 155 160

Tyr Ala Lys Asn Gly Glu Leu Leu Lys Tyr Ile Arg Lys Ile Gly Ser
165 170 175

Phe Asp Glu Thr Cys Thr Arg Phe Tyr Thr Ala Glu Ile Val Ser Ala
180 185 190

Leu Glu Tyr Leu His Gly Lys Gly Ile Ile His Arg Asp Leu Lys Pro
195 200 205

Glu Asn Ile Leu Leu Asn Glu Asp Met His Ile Gln Ile Thr Asp Phe
210 215 220

Gly Thr Ala Lys Val Leu Ser Pro Glu Ser Lys Gln Val Cys
225 230 235

<210> 155

<211> 73

<212> PRT

<213> Homo sapiens

<400> 155

Met Ser Asp Val Thr Ile Val Lys Glu Gly Trp Val Gln Lys Arg Gly
1 5 10 15

Glu Tyr Ile Lys Asn Trp Arg Pro Arg Tyr Phe Leu Leu Lys Thr Asp
20 25 30

Gly Ser Phe Ile Gly Tyr Lys Glu Lys Pro Gln Asp Val Asp Leu Pro
35 40 45

Tyr Pro Leu Asn Asn Phe Ser Val Ala Ser Ser Val Met Phe Arg Tyr
50 55 60

Leu Gln Asn Leu Thr Leu Asn Gln Val
65 70

<210> 156

<211> 213

<212> PRT

<213> Homo sapiens

<400> 156

Met Ser Asp Val Thr Ile Val Lys Glu Gly Trp Val Gln Lys Arg Gly
1 5 10 15

Glu Tyr Ile Lys Asn Trp Arg Pro Arg Tyr Phe Leu Leu Lys Thr Asp
20 25 30

Gly Ser Phe Ile Gly Tyr Lys Glu Lys Pro Gln Asp Val Asp Leu Pro
35 40 45

Tyr Pro Leu Asn Asn Phe Ser Val Ala Lys Cys Gln Leu Met Lys Thr
50 55 60

Glu Arg Pro Lys Pro Asn Thr Phe Ile Ile Arg Cys Leu Gln Trp Thr
65 70 75 80

Thr Val Ile Glu Arg Thr Phe His Val Asp Thr Pro Glu Glu Arg Glu
85 90 95

Glu Trp Thr Glu Ala Ile Gln Ala Val Ala Asp Arg Leu Gln Arg Gln
100 105 110

Glu Glu Glu Arg Met Asn Cys Ser Pro Thr Ser Gln Ile Asp Asn Ile
115 120 125

Gly Glu Glu Glu Met Asp Ala Ser Thr Thr His His Lys Arg Lys Thr
130 135 140

Met Asn Asp Phe Asp Tyr Leu Lys Leu Leu Gly Lys Gly Thr Phe Gly
145 150 155 160

Lys Val Ile Leu Val Arg Glu Lys Ala Ser Gly Lys Tyr Tyr Ala Met
165 170 175

Lys Ile Leu Lys Lys Glu Val Ile Ile Ala Lys Val Thr Asp Leu Leu
180 185 190

Lys Leu Ile Thr Lys Phe Leu Phe Ala Val Cys Met Cys Leu Trp Ala
195 200 205

His Glu Phe Thr Cys
210

<210> 157

<211> 352

<212> PRT

<213> Homo sapiens

<400> 157

Met Gly Gly Lys Pro Ala Asn Arg Met Met Pro Tyr Pro Phe Pro Ser
1 5 10 15

Gly Thr Trp Lys Val Lys Trp Val Ala Ser Arg Asn Ala Phe Lys Pro
20 25 30

Arg Ile Gly Ile Leu Ile Lys Thr Leu Ile Tyr Ser Ser Gln Phe Pro
35 40 45

Leu Gly Asn Leu Glu Lys Ile Ser Gln Leu Leu Ser Lys Ser Ala Gln
50 55 60

Cys Pro Leu Arg Val His Tyr Leu Ser Ser Gln Tyr Gly Asp Glu Arg
65 70 75 80

Cys Phe Met Phe Val Leu Ile Ser Pro Thr Lys Ser Val Ile Ile Thr
85 90 95

Ile Leu Ser Leu Leu Phe Thr Leu Gln Leu Phe Phe His Leu Ser Arg
100 105 110

Glu Arg Val Phe Ser Glu Asp Arg Thr Arg Phe Tyr Gly Ala Glu Ile
115 120 125

Val Ser Ala Leu Asp Tyr Leu His Ser Gly Lys Ile Val Tyr Arg Asp
130 135 140

Leu Lys Leu Glu Asn Leu Met Leu Asp Lys Asp Gly His Ile Lys Ile
145 150 155 160

Thr Asp Phe Gly Leu Cys Lys Glu Gly Ile Thr Asp Ala Ala Thr Met
165 170 175

Lys Thr Phe Cys Gly Thr Pro Glu Tyr Leu Ala Pro Glu Val Leu Glu
180 185 190

Asp Asn Asp Tyr Gly Arg Ala Val Asp Trp Trp Gly Leu Gly Val Val
195 200 205

Met Tyr Glu Met Met Cys Gly Arg Leu Pro Phe Tyr Asn Gln Asp His
210 215 220

Glu Lys Leu Phe Glu Leu Ile Leu Met Glu Asp Ile Lys Phe Pro Arg
225 230 235 240

Thr Leu Ser Ser Asp Ala Lys Ser Leu Leu Ser Gly Leu Leu Ile Lys
245 250 255

Asp Pro Asn Lys Arg Leu Gly Gly Pro Asp Asp Ala Lys Glu Ile
260 265 270

Met Arg His Ser Phe Phe Ser Gly Val Asn Trp Gln Asp Val Tyr Asp
275 280 285

Lys Lys Leu Val Pro Pro Phe Lys Pro Gln Val Thr Ser Glu Thr Asp
290 295 300

Thr Arg Tyr Phe Asp Glu Glu Phe Thr Ala Gln Thr Ile Thr Ile Thr
305 310 315 320

Pro Pro Glu Lys Tyr Asp Glu Asp Gly Met Asp Cys Met Asp Asn Glu
325 330 335

Arg Arg Pro His Phe Pro Gln Phe Ser Tyr Ser Ala Ser Gly Arg Glu
340 345 350

<210> 158
<211> 132
<212> PRT
<213> Homo sapiens

<400> 158

Met Glu Leu Leu Arg Thr Ile Thr Tyr Gln Pro Ala Ala Ser Thr Lys
1 5 10 15

Met Cys Glu Gln Ala Leu Gly Lys Gly Cys Gly Asn Ser Lys Lys
20 25 30

Lys Arg Pro Pro Gln Pro Pro Glu Glu Ser Gln Pro Pro Gln Ser Gln
35 40 45

Ala Gln Val Pro Pro Ala Ala Pro His His His His His Ser His
50 55 60

Ser Gly Pro Glu Ile Ser Arg Ile Ile Val Asp Pro Thr Thr Gly Lys
65 70 75 80

Arg Tyr Cys Arg Gly Lys Val Leu Gly Lys Gly Gly Phe Ala Lys Cys
85 90 95

Tyr Glu Met Thr Asp Leu Thr Asn Asn Lys Val Tyr Ala Ala Lys Ile
100 105 110

Ile Pro His Ser Arg Val Ala Lys Pro His Gln Arg Glu Lys Val Cys
115 120 125

Met Thr Leu Glu
130

<210> 159
<211> 192
<212> PRT
<213> Homo sapiens

<400> 159

Met Glu Leu Leu Arg Thr Ile Thr Tyr Gln Pro Ala Ala Ser Thr Lys
1 5 10 15

Met Cys Glu Gln Ala Leu Gly Lys Gly Cys Gly Asn Ser Lys Lys
20 25 30

Lys Arg Pro Pro Gln Pro Pro Glu Glu Ser Gln Pro Pro Gln Ser Gln
35 40 45

Ala Gln Val Pro Pro Ala Ala Pro His His His His His Ser His
50 55 60

Ser Gly Pro Glu Ile Ser Arg Ile Ile Val Asp Pro Thr Thr Gly Lys

65 70 75 80

Arg Tyr Cys Arg Gly Lys Val Leu Gly Lys Gly Gly Phe Ala Lys Cys
85 90 95

Tyr Glu Met Thr Asp Leu Thr Asn Asn Lys Val Tyr Ala Ala Lys Ile
100 105 110

Ile Pro His Ser Arg Val Ala Lys Pro His Gln Arg Glu Lys Ile Asp
115 120 125

Lys Glu Ile Glu Leu His Arg Ile Leu His His Lys His Val Val Gln
130 135 140

Phe Tyr His Tyr Phe Glu Asp Lys Glu Asn Ile Tyr Ile Leu Leu Glu
145 150 155 160

Tyr Cys Ser Arg Arg Val Ser Val Asn Ser Tyr Leu Arg Thr Phe Ala
165 170 175

Tyr Pro Glu Leu Thr Trp Tyr Ser Lys Ser Ile Leu Ser Gly Ile Thr
180 185 190

<210> 160
<211> 207
<212> PRT
<213> Homo sapiens

<400> 160

Met Glu Leu Leu Arg Thr Ile Thr Tyr Gln Pro Ala Ala Ser Thr Lys
1 5 10 15

Met Cys Glu Gln Ala Leu Gly Lys Gly Cys Gly Gly Asn Ser Lys Lys
20 25 30

Lys Arg Pro Pro Gln Pro Pro Glu Glu Ser Gln Pro Pro Gln Ser Gln
35 40 45

Ala Gln Val Pro Pro Ala Ala Pro His His His His His Ser His
50 55 60

Ser Gly Pro Glu Ile Ser Arg Ile Ile Val Asp Pro Thr Thr Gly Lys
65 70 75 80

Arg Tyr Cys Arg Gly Lys Val Leu Gly Lys Gly Gly Phe Ala Lys Cys
85 90 95

Tyr Glu Met Thr Asp Leu Thr Asn Asn Lys Val Tyr Ala Ala Lys Ile
100 105 110

Ile Pro His Ser Arg Val Ala Lys Pro His Gln Arg Glu Lys Ile Asp
115 120 125

Lys Glu Ile Glu Leu His Arg Ile Leu His His Lys His Val Val Gln
130 135 140

Phe Tyr His Tyr Phe Glu Asp Lys Glu Asn Ile Tyr Ile Leu Leu Glu
145 150 155 160

Tyr Cys Ser Arg Arg Leu Gln Gly Ser Gln Lys Asn Asp Leu Glu Tyr
165 170 175

Val Glu Glu Asp Gly His Val Val Val Arg Lys Gln Phe Pro Cys Gly
180 185 190

Leu Leu Asp Trp Val Glu Pro Glu Gln Ala Lys Ala Tyr Ser Ser
195 200 205

<210> 161
<211> 337
<212> PRT
<213> Homo sapiens

<400> 161

Met Ser Asp Lys Asp Leu Arg Thr Ala Ala Ala Gly Gly Gly His Leu
1 5 10 15

Val Ala Ile Leu Thr Val Phe Ile Pro Gln Lys Asp Leu Val Glu Glu
20 25 30

Glu Ala Glu Glu Ala Gly Val Ala Leu Arg Ser Thr Gln Ser Thr Leu
35 40 45

Gln Ala Gly Leu Ala Ala Asp Ala Trp Ala Ala Pro Ile Ala Met Gln
50 55 60

Ile Tyr Lys Lys His Leu Asp Pro Arg Pro Gly Pro Cys His Leu Ser
65 70 75 80

Trp Ala Trp Ala Trp Ala Ser Trp Pro Ala Ala Ala Cys Thr Ala Gly
85 90 95

Pro Lys Gly Arg Pro Pro Met Thr Gln Val Tyr Glu Arg Leu Glu Lys
100 105 110

Leu Gln Ala Val Val Ala Gly Val Pro Gly His Leu Glu Ala Ala Ser
115 120 125

Cys Ile Pro Phe Pro Gln Glu Asn Ser Tyr Val Ser Ser Thr Gly Arg
130 135 140

Ala Ser Ala Gln Ala Ala Glu Gln Leu Gln Arg Gly Pro Asn Gln Pro
145 150 155 160

Val Glu Ser Asp Glu Ser Leu Gly Gly Leu Ser Ala Ala Leu Arg Ser
165 170 175

Trp His Leu Thr Pro Ser Cys Pro Leu Asp Pro Ala Pro Leu Arg Glu
180 185 190

Ala Gly Cys Pro Gln Gly Asp Thr Ala Gly Glu Ser Ser Trp Gly Ser
195 200 205

Gly Pro Gly Ser Arg Pro Thr Ala Val Glu Gly Leu Ala Leu Gly Ser
210 215 220

Ser Ala Ser Ser Ser Glu Pro Pro Gln Ile Ile Asn Pro Ala
225 230 235 240

Arg Gln Lys Met Val Gln Lys Leu Ala Leu Tyr Glu Asp Gly Ala Leu
245 250 255

Asp Ser Leu Gln Leu Leu Ser Ser Ser Leu Pro Gly Leu Gly Leu
260 265 270

Glu Gln Asp Arg Gln Gly Pro Lys Lys Val Met Asn Phe Arg Ala Asp
275 280 285

Val Phe Thr Trp Ala Asp Pro Pro Asn Pro Glu Val Lys Val Leu Met
290 295 300

Val Arg Ser Ser His Gly Ala Arg Val Leu Ser Thr Leu Pro Ala Val
305 310 315 320

Gly Val Gly Ala His Ala Arg Trp Gly Glu Lys Glu Val Ala Leu Leu
325 330 335

Phe

<210> 162
<211> 122
<212> PRT
<213> Homo sapiens

<400> 162

Met Gly His Ala Leu Cys Val Cys Ser Arg Gly Thr Val Ile Ile Asp
1 5 10 15

Asn Lys Arg Tyr Leu Phe Ile Gln Lys Leu Gly Glu Gly Phe Ser
20 25 30

Tyr Val Asp Leu Val Glu Gly Leu His Asp Gly His Phe Tyr Ala Leu
35 40 45

Lys Arg Ile Leu Cys His Glu Gln Gln Asp Arg Glu Glu Ala Gln Arg
50 55 60

Glu Ala Asp Met His Arg Leu Phe Asn His Pro Asn Ile Leu Arg Leu
65 70 75 80

Val Ala Tyr Cys Leu Arg Glu Arg Gly Ala Lys His Glu Ala Trp Leu
85 90 95

Leu Leu Pro Phe Phe Lys Val Arg Lys Thr Pro Val Tyr Gly Gly Gly
100 105 110

Cys Ser Arg Ala Thr Tyr Ser Arg Ala Val

115

120

<210> 163
<211> 842
<212> PRT
<213> Homo sapiens

<400> 163

Met Glu Arg Ala Ile Ser Pro Gly Leu Leu Val Arg Ala Leu Leu Leu
1 5 10 15

Leu Leu Leu Leu Gly Leu Ala Ala Arg Thr Val Ala Ala Gly Arg Ala
20 25 30

Arg Gly Leu Pro Ala Pro Thr Ala Glu Ala Ala Phe Gly Leu Gly Ala
35 40 45

Ala Ala Ala Pro Thr Ser Ala Thr Arg Val Pro Ala Ala Gly Ala Val
50 55 60

Ala Ala Ala Glu Val Thr Val Glu Asp Ala Glu Ala Leu Pro Ala Ala
65 70 75 80

Ala Gly Glu Gln Glu Pro Arg Gly Pro Glu Pro Asp Asp Glu Thr Glu
85 90 95

Leu Arg Pro Arg Gly Arg Ser Leu Val Ile Ile Ser Thr Leu Asp Gly
100 105 110

Arg Ile Ala Ala Leu Asp Pro Glu Asn His Gly Lys Lys Gln Trp Asp
115 120 125

Leu Asp Val Gly Ser Gly Ser Leu Val Ser Ser Ser Leu Ser Lys Pro
130 135 140

Glu Val Phe Gly Asn Lys Met Ile Ile Pro Ser Leu Asp Gly Ala Leu
145 150 155 160

Phe Gln Trp Asp Arg Asp Arg Glu Ser Met Glu Thr Val Pro Phe Thr
165 170 175

Val Glu Ser Leu Leu Glu Ser Ser Tyr Lys Phe Gly Asp Asp Val Val
180 185 190

Leu Val Gly Gly Lys Ser Leu Thr Thr Tyr Gly Leu Ser Ala Tyr Ser
195 200 205

Gly Lys Val Arg Tyr Ile Cys Ser Ala Leu Gly Cys Arg Gln Trp Asp
210 215 220

Ser Asp Glu Met Glu Gln Glu Glu Asp Ile Leu Leu Leu Gln Arg Thr
225 230 235 240

Gln Lys Thr Val Arg Ala Val Gly Pro Arg Ser Gly Asn Glu Lys Trp
245 250 255

Asn Phe Ser Val Gly His Phe Glu Leu Arg Tyr Ile Pro Asp Met Glu
260 265 270

Thr Arg Ala Gly Phe Ile Glu Ser Thr Phe Lys Pro Asn Glu Asn Thr
275 280 285

Glu Glu Ser Lys Ile Ile Ser Asp Val Glu Glu Gln Glu Ala Ala Ile
290 295 300

Met Asp Ile Val Ile Lys Val Ser Val Ala Asp Trp Lys Val Met Ala
305 310 315 320

Phe Ser Lys Lys Gly Gly His Leu Glu Trp Glu Tyr Gln Phe Cys Thr
325 330 335

Pro Ile Ala Ser Ala Trp Leu Leu Lys Asp Gly Lys Val Ile Pro Ile
340 345 350

Ser Leu Phe Asp Asp Thr Ser Tyr Thr Ser Asn Asp Asp Val Leu Glu
355 360 365

Asp Glu Glu Asp Ile Val Glu Ala Ala Arg Gly Ala Thr Glu Asn Ser
370 375 380

Val Tyr Leu Gly Met Tyr Arg Gly Gln Leu Tyr Leu Gln Ser Ser Val
385 390 395 400

Arg Ile Ser Glu Lys Phe Pro Ser Ser Pro Lys Ala Leu Glu Ser Val
405 410 415

Thr Asn Glu Asn Ala Ile Ile Pro Leu Pro Thr Ile Lys Trp Lys Pro
420 425 430

Leu Ile His Ser Pro Ser Arg Thr Pro Val Leu Val Gly Ser Asp Glu
435 440 445

Phe Asp Lys Cys Leu Ser Asn Asp Lys Phe Ser His Glu Glu Tyr Ser
450 455 460

Asn Gly Ala Leu Ser Ile Leu Gln Tyr Pro Tyr Asp Asn Gly Tyr Tyr
465 470 475 480

Leu Pro Tyr Tyr Lys Arg Glu Arg Asn Lys Arg Ser Thr Gln Ile Thr
485 490 495

Val Arg Phe Leu Asp Asn Pro His Tyr Asn Lys Asn Ile Arg Lys Lys
500 505 510

Asp Pro Val Leu Leu His Trp Trp Lys Glu Ile Val Ala Thr Ile
515 520 525

Leu Phe Cys Ile Ile Ala Thr Thr Phe Ile Val Arg Arg Leu Phe His
530 535 540

Pro His Pro His Arg Gln Arg Lys Glu Ser Glu Thr Gln Cys Gln Thr
545 550 555 560

Glu Asn Lys Tyr Asp Ser Val Ser Gly Glu Ala Asn Asp Ser Ser Trp
565 570 575

Asn Asp Ile Lys Asn Ser Gly Tyr Ile Ser Arg Tyr Leu Thr Asp Phe
580 585 590

Glu Pro Ile Gln Cys Leu Gly Arg Gly Phe Gly Val Val Phe Glu
595 600 605

Ala Lys Asn Lys Val Asp Asp Cys Asn Tyr Ala Ile Lys Arg Ile Arg
610 615 620

Leu Pro Asn Arg Glu Leu Ala Arg Glu Lys Val Met Arg Glu Val Lys
625 630 635 640

Ala Leu Ala Lys Leu Glu His Pro Gly Ile Val Arg Tyr Phe Asn Ala
645 650 655

Trp Leu Glu Ala Pro Pro Glu Lys Trp Gln Glu Lys Met Asp Glu Ile
660 665 670

Trp Leu Lys Asp Glu Ser Thr Asp Trp Pro Leu Ser Ser Pro Ser Pro
675 680 685

Met Asp Ala Pro Ser Val Lys Ile Arg Arg Met Asp Pro Phe Ser Thr
690 695 700

Lys Glu His Ile Glu Ile Ile Ala Pro Ser Pro Gln Arg Ser Arg Ser
705 710 715 720

Phe Ser Val Gly Ile Ser Cys Asp Gln Thr Ser Ser Ser Glu Ser Gln
725 730 735

Phe Ser Pro Leu Glu Phe Ser Gly Met Asp His Glu Asp Ile Ser Glu
740 745 750

Ser Val Asp Ala Ala Tyr Asn Leu Gln Asp Ser Cys Leu Thr Asp Cys
755 760 765

Asp Val Glu Asp Gly Thr Met Asp Gly Asn Asp Glu Gly His Ser Phe
770 775 780

Glu Leu Cys Pro Ser Glu Ala Ser Pro Tyr Val Arg Ser Arg Glu Arg
785 790 795 800

Thr Ser Ser Ser Ile Val Phe Glu Asp Ser Gly Cys Asp Asn Ala Ser
805 810 815

Ser Lys Glu Glu Pro Lys Thr Asn Arg Leu His Ile Gly Asn His Cys
820 825 830

Ala Asn Lys Leu Thr Val Thr Val Leu Phe
835 840

<210> 164
<211> 743
<212> PRT

<213> Homo sapiens

<400> 164

Met Gly Ser Arg Ala Gln Lys Ser Ala Gly Asn Ala Glu Leu Trp Glu
1 5 10 15

Pro Leu Pro Glu Gly Arg Pro Arg Pro Ala Gly Thr Ser Ser Ala Val
20 25 30

Ser Ala Trp Ala Ser Leu Lys Leu Cys Leu Arg Gly Gly Ser Gly Arg
35 40 45

Arg Gln Arg Leu Gly Gly Arg Met Gln Pro Glu Glu Gly His Arg
50 55 60

Leu Ala Ala Gly Ala Ala Val Arg Gly Ala Ala Ala Thr Val Leu Leu
65 70 75 80

Arg Leu Arg Asp Asp Leu Asn Val Thr Arg Leu Ser His Phe Glu Tyr
85 90 95

Val Lys Asn Glu Asp Leu Glu Lys Ile Gly Met Gly Arg Pro Gly Gln
100 105 110

Arg Arg Leu Trp Glu Ala Val Lys Arg Arg Lys Ala Leu Cys Lys Arg
115 120 125

Lys Ser Trp Met Asn Lys Val Phe Ser Gly Lys Arg Leu Glu Ala Glu
130 135 140

Phe Pro Pro His His Ser Gln Ser Thr Phe Arg Lys Thr Ser Pro Ala
145 150 155 160

Pro Gly Gly Pro Ala Gly Glu Gly Pro Leu Gln Ser Leu Thr Cys Leu
165 170 175

Ile Gly Glu Lys Asp Leu Arg Leu Leu Glu Lys Leu Gly Asp Gly Ser
180 185 190

Phe Gly Val Val Arg Arg Gly Glu Trp Asp Ala Pro Ser Gly Lys Thr
195 200 205

Val Ser Pro Pro Gln Pro Ala Phe Phe Thr Gln Lys Pro Thr Tyr Asp
210 215 220

Pro Val Ser Glu Asp Gln Asp Pro Leu Ser Ser Asp Phe Lys Arg Leu
225 230 235 240

Gly Leu Arg Lys Pro Gly Leu Pro Arg Gly Leu Trp Leu Ala Lys Pro
245 250 255

Ser Ala Arg Val Pro Gly Thr Lys Ala Ser Arg Gly Ser Gly Ala Glu
260 265 270

Val Thr Leu Ile Asp Phe Gly Glu Glu Pro Val Val Pro Ala Leu Arg
275 280 285

Pro Cys Ala Pro Ser Leu Ala Gln Leu Ala Met Asp Ala Cys Ser Leu
290 295 300

Leu Asp Glu Thr Pro Pro Gln Ser Pro Thr Arg Ala Leu Pro Arg Pro
305 310 315 320

Leu His Pro Thr Pro Val Val Asp Trp Asp Ala Arg Pro Leu Pro Pro
325 330 335

Pro Pro Ala Tyr Asp Asp Val Ala Gln Asp Glu Asp Asp Phe Glu Ile
340 345 350

Cys Ser Ile Asn Ser Thr Leu Val Gly Ala Gly Val Pro Ala Gly Pro
355 360 365

Ser Gln Gly Gln Thr Asn Tyr Ala Phe Val Pro Glu Gln Ala Arg Pro
370 375 380

Pro Pro Pro Leu Glu Asp Asn Leu Phe Leu Pro Pro Gln Gly Gly
385 390 395 400

Lys Pro Pro Ser Ser Ala Gln Thr Ala Glu Ile Phe Gln Ala Leu Gln
405 410 415

Gln Glu Cys Met Arg Gln Leu Gln Ala Pro Ala Gly Ser Pro Ala Pro
420 425 430

Ser Pro Ser Pro Gly Gly Asp Asp Lys Pro Gln Val Pro Pro Arg Val
435 440 445

Pro Ile Pro Pro Arg Pro Thr Arg Pro His Val Gln Leu Ser Pro Ala
450 455 460

Pro Pro Gly Glu Glu Glu Thr Ser Gln Trp Pro Gly Pro Ala Ser Pro
465 470 475 480

Pro Arg Val Pro Pro Arg Glu Pro Leu Ser Pro Gln Gly Ser Arg Thr
485 490 495

Pro Ser Pro Leu Val Pro Pro Gly Ser Ser Pro Leu Pro Pro Arg Leu
500 505 510

Ser Ser Ser Pro Gly Lys Thr Met Pro Thr Thr Gln Ser Phe Ala Ser
515 520 525

Asp Pro Lys Tyr Ala Thr Pro Gln Val Ile Gln Ala Pro Gly Pro Arg
530 535 540

Ala Gly Pro Cys Ile Leu Pro Ile Val Arg Asp Gly Lys Lys Val Ser
545 550 555 560

Ser Thr His Tyr Tyr Leu Leu Pro Glu Arg Pro Ser Tyr Leu Glu Arg
565 570 575

Tyr Gln Arg Phe Leu Arg Glu Ala Gln Ser Pro Glu Glu Pro Thr Pro
580 585 590

Leu Pro Val Pro Leu Leu Leu Pro Pro Pro Ser Thr Pro Ala Pro Ala
595 600 605

Ala Pro Thr Ala Thr Val Arg Pro Met Pro Gln Ala Ala Leu Asp Pro
610 615 620

Lys Ala Asn Phe Ser Thr Asn Asn Ser Asn Pro Gly Ala Arg Pro Pro
625 630 635 640

Pro Pro Arg Ala Thr Ala Arg Leu Pro Gln Arg Gly Cys Pro Gly Asp
645 650 655

Gly Pro Glu Ala Gly Arg Pro Ala Asp Lys Ile Gln Met Ala Met Val.
660 665 670

His Gly Val Thr Thr Glu Glu Cys Gln Ala Ala Leu Gln Cys His Gly
675 680 685

Trp Ser Val Gln Arg Ala Cys Pro Val Ser Glu Gly Gly Ala Ala Leu
690 695 700

Arg Ala Gly Ser Ala Ala Gln Arg Glu Cys His Lys Val Leu Glu Met
705 710 715 720

Phe Asp Trp Asn Leu Glu Gln Ala Gly Cys His Leu Leu Gly Ser Trp
725 730 735

Gly Pro Ala His His Lys Arg
740

<210> 165
<211> 604
<212> PRT
<213> Homo sapiens

<400> 165

Met Ala Ser Asn Pro Glu Arg Gly Glu Ile Leu Leu Thr Glu Leu Gln
1 5 10 15

Gly Asp Ser Arg Ser Leu Pro Phe Ser Glu Asn Val Ser Ala Val Gln
20 25 30

Lys Leu Asp Phe Ser Asp Thr Met Val Gln Gln Lys Leu Asp Asp Ile
35 40 45

Lys Asp Arg Ile Lys Arg Glu Ile Arg Lys Glu Leu Lys Ile Lys Glu
50 55 60

Gly Ala Glu Asn Leu Arg Lys Val Thr Thr Asp Lys Lys Ser Leu Ala
65 70 75 80

Tyr Val Asp Asn Ile Leu Lys Ser Asn Lys Lys Leu Glu Leu
85 90 95

His His Lys Leu Gln Glu Leu Asn Ala His Ile Val Val Ser Asp Pro

100 105 110

Glu Asp Ile Thr Asp Cys Pro Arg Thr Pro Asp Thr Pro Asn Asn Asp
115 120 125

Pro Arg Cys Ser Thr Ser Asn Asn Arg Leu Lys Ala Leu Gln Lys Gln
130 135 140

Leu Asp Ile Glu Leu Lys Val Lys Gln Gly Ala Glu Asn Met Ile Gln
145 150 155 160

Met Tyr Ser Asn Gly Ser Ser Lys Asp Arg Lys Leu His Gly Thr Ala
165 170 175

Gln Gln Leu Leu Gln Asp Ser Lys Thr Lys Ile Glu Val Ile Arg Met
180 185 190

Gln Ile Leu Gln Ala Val Gln Thr Asn Glu Leu Ala Phe Asp Asn Ala
195 200 205

Lys Pro Val Ile Ser Pro Leu Glu Leu Arg Met Glu Glu Leu Arg His
210 215 220

His Phe Arg Ile Glu Phe Ala Val Ala Glu Gly Ala Lys Asn Val Met
225 230 235 240

Lys Leu Leu Gly Ser Gly Lys Val Thr Asp Arg Lys Ala Leu Ser Glu
245 250 255

Ala Gln Ala Arg Phe Asn Glu Ser Ser Gln Lys Leu Asp Leu Leu Lys
260 265 270

Tyr Ser Leu Glu Gln Arg Leu Asn Glu Val Pro Lys Asn His Pro Lys
275 280 285

Ser Arg Ile Ile Ile Glu Glu Leu Ser Leu Val Ala Ala Ser Pro Thr
290 295 300

Leu Ser Pro Arg Gln Ser Met Ile Ser Thr Gln Asn Gln Tyr Ser Thr
305 310 315 320

Leu Ser Lys Pro Ala Ala Leu Thr Gly Thr Leu Glu Val Arg Leu Met
325 330 335

Gly Cys Gln Asp Ile Leu Glu Asn Val Pro Gly Arg Ser Lys Ala Thr
340 345 350

Ser Val Ala Leu Pro Gly Trp Ser Pro Ser Glu Thr Arg Ser Ser Phe
355 360 365

Met Ser Arg Thr Ser Lys Ser Lys Ser Gly Ser Ser Arg Asn Leu Leu
370 375 380

Lys Thr Asp Asp Leu Ser Asn Asp Val Cys Ala Val Leu Lys Leu Asp
385 390 395 400

Asn Thr Val Val Gly Gln Thr Ser Trp Lys Pro Ile Ser Asn Gln Ser

405

410

415

Trp Asp Gln Lys Phe Thr Leu Glu Leu Asp Arg Ser Arg Glu Leu Glu
420 425 430

Ile Ser Val Tyr Trp Arg Asp Trp Arg Ser Leu Cys Ala Val Lys Phe
435 440 445

Leu Arg Leu Glu Asp Phe Leu Asp Asn Gln Arg His Gly Met Cys Leu
450 455 460

Tyr Leu Glu Pro Gln Gly Thr Leu Phe Ala Glu Val Thr Phe Phe Asn
465 470 475 480

Pro Val Ile Glu Arg Arg Pro Lys Leu Gln Arg Gln Lys Lys Ile Phe
485 490 495

Ser Lys Gln Gln Gly Lys Thr Phe Leu Arg Ala Pro Gln Met Asn Ile
500 505 510

Asn Ile Ala Thr Trp Gly Arg Leu Val Arg Arg Ala Ile Pro Thr Val
515 520 525

Asn His Ser Gly Thr Phe Ser Pro Gln Ala Pro Val Pro Thr Thr Val
530 535 540

Pro Val Val Asp Val Arg Ile Pro Gln Leu Ala Pro Pro Ala Arg Tyr
545 550 555 560

Val Ser Glu Ile Leu Ser Ile Ser Tyr Thr Lys Leu Leu Gly His Ser
565 570 575

Tyr Val Leu Ile Ile Ala Gly Val Leu Ser Leu Ala Phe Phe Pro Ser
580 585 590

Ser Ile Leu Lys Val Val Phe Cys Leu Leu Lys Lys
595 600

<210> 166

<211> 613

<212> PRT

<213> Homo sapiens

<400> 166

Met Ala Ser Asn Pro Glu Arg Gly Glu Ile Leu Leu Thr Glu Leu Gln
1 5 10 15

Gly Asp Ser Arg Ser Leu Pro Phe Ser Glu Asn Val Ser Ala Val Gln
20 25 30

Lys Leu Asp Phe Ser Asp Thr Met Val Gln Gln Lys Leu Asp Asp Ile
35 40 45

Lys Asp Arg Ile Lys Arg Glu Ile Arg Lys Glu Leu Lys Ile Lys Glu
50 55 60

Gly Ala Glu Asn Leu Arg Lys Val Thr Thr Asp Lys Lys Ser Leu Ala
65 70 75 80

Tyr Val Asp Asn Ile Leu Lys Lys Ser Asn Lys Lys Leu Glu Glu Leu
85 90 95

His His Lys Leu Gln Glu Leu Asn Ala His Ile Val Val Ser Asp Pro
100 105 110

Glu Asp Ile Thr Asp Cys Pro Arg Thr Pro Asp Thr Pro Asn Asn Asp
115 120 125

Pro Arg Cys Ser Thr Ser Asn Asn Arg Leu Lys Ala Leu Gln Lys Gln
130 135 140

Leu Asp Ile Glu Leu Lys Val Lys Gln Gly Ala Glu Asn Met Ile Gln
145 150 155 160

Met Tyr Ser Asn Gly Ser Ser Lys Asp Arg Lys Leu His Gly Thr Ala
165 170 175

Gln Gln Leu Leu Gln Asp Ser Lys Thr Lys Ile Glu Val Ile Arg Met
180 185 190

Gln Ile Leu Gln Ala Val Gln Thr Asn Glu Leu Ala Phe Asp Asn Ala
195 200 205

Lys Pro Val Ile Ser Pro Leu Glu Leu Arg Met Glu Glu Leu Arg His
210 215 220

His Phe Arg Ile Glu Phe Ala Val Ala Glu Gly Ala Lys Asn Val Met
225 230 235 240

Lys Leu Leu Gly Ser Gly Lys Val Thr Asp Arg Lys Ala Leu Ser Glu
245 250 255

Ala Gln Ala Arg Phe Asn Glu Ser Ser Gln Lys Leu Asp Leu Leu Lys
260 265 270

Tyr Ser Leu Glu Gln Arg Leu Asn Glu Val Pro Lys Asn His Pro Lys
275 280 285

Ser Arg Ile Ile Ile Glu Glu Leu Ser Leu Val Ala Ala Ser Pro Thr
290 295 300

Leu Ser Pro Arg Gln Ser Met Ile Ser Thr Gln Asn Gln Tyr Ser Thr
305 310 315 320

Leu Ser Lys Pro Ala Ala Leu Thr Gly Thr Leu Glu Val Arg Leu Met
325 330 335

Gly Cys Gln Asp Ile Leu Glu Asn Val Pro Gly Arg Ser Lys Ala Thr
340 345 350

Ser Val Ala Leu Pro Gly Trp Ser Pro Ser Glu Thr Arg Ser Ser Phe
355 360 365

Met Ser Arg Thr Ser Lys Ser Lys Ser Gly Ser Ser Arg Asn Leu Leu
370 375 380

Lys Thr Asp Asp Leu Ser Asn Asp Val Cys Ala Val Leu Lys Leu Asp
385 390 395 400

Asn Thr Val Val Gly Gln Thr Ser Trp Lys Pro Ile Ser Asn Gln Ser
405 410 415

Trp Asp Gln Lys Phe Thr Leu Glu Leu Asp Arg Ser Arg Glu Leu Glu
420 425 430

Ile Ser Val Tyr Trp Arg Asp Trp Arg Ser Leu Cys Ala Val Lys Phe
435 440 445

Leu Arg Leu Glu Asp Phe Leu Asp Asn Gln Arg His Gly Met Cys Leu
450 455 460

Tyr Leu Glu Pro Gln Gly Thr Leu Phe Ala Glu Val Thr Phe Phe Asn
465 470 475 480

Pro Val Ile Glu Arg Arg Pro Lys Leu Gln Arg Gln Lys Lys Ile Phe
485 490 495

Ser Lys Gln Gln Gly Lys Thr Phe Leu Arg Ala Pro Gln Met Asn Ile
500 505 510

Asn Ile Ala Thr Trp Gly Arg Leu Val Arg Arg Ala Ile Pro Thr Val
515 520 525

Asn His Ser Gly Thr Phe Ser Pro Gln Ala Pro Val Pro Thr Thr Val
530 535 540

Pro Val Val Asp Val Arg Ile Pro Gln Leu Ala Pro Pro Ala Ser Asp
545 550 555 560

Ser Thr Val Thr Lys Leu Asp Phe Asp Leu Glu Pro Glu Pro Pro Pro
565 570 575

Ala Pro Pro Arg Ala Ser Ser Leu Gly Glu Ile Asp Glu Ser Ser Glu
580 585 590

Leu Arg Val Leu Asp Ile Pro Gly Gln Ala Ser His Phe Lys Pro Cys
595 600 605

Ile Ile Pro Leu His
610

<210> 167
<211> 133
<212> PRT
<213> Homo sapiens

<400> 167

Met Val Ser Ser Gln Lys Leu Glu Lys Pro Ile Glu Met Gly Ser Ser
1 5 10 15

Glu Pro Leu Pro Ile Ala Asp Gly Asp Arg Arg Arg Lys Lys Lys Arg
20 25 30

Arg Gly Arg Ala Thr Asp Ser Leu Pro Gly Lys Phe Glu Asp Met Tyr
35 40 45

Lys Leu Thr Ser Glu Leu Leu Gly Glu Gly Ala Tyr Ala Lys Val Gln
50 55 60

Gly Ala Val Ser Leu Gln Asn Gly Lys Glu Tyr Ala Val Lys Ile Ile
65 70 75 80

Glu Lys Gln Ala Gly His Ser Arg Ser Arg Val Phe Arg Glu Val Glu
85 90 95

Thr Leu Tyr Gln Cys Gln Gly Asn Lys Asn Ile Leu Glu Leu Ile Glu
100 105 110

Phe Phe Glu Asp Asp Thr Arg Phe Tyr Leu Val Phe Glu Lys Leu Gln
115 120 125

Gly Gly Thr Tyr Arg
130

<210> 168
<211> 153
<212> PRT
<213> Homo sapiens

<400> 168

Met Leu Gln Val Gly Val Leu Arg Asp Arg Ser Pro Ala Gly Ala Ser
1 5 10 15

Glu Gly Phe His Val Arg Gly Arg Trp Arg Thr Glu Asp Cys His Leu
20 25 30

Arg Thr Lys Ala Ile Glu Thr Leu Arg Val Ala Gly Arg His Gln Leu
35 40 45

Pro Asp Arg Ser Phe Ile Ser Phe Gly Ile Ser Ser Leu Gln Met Val
50 55 60

Ser Ser Gln Lys Leu Glu Lys Pro Ile Glu Met Gly Ser Ser Glu Pro
65 70 75 80

Leu Pro Ile Ala Asp Gly Asp Arg Arg Arg Lys Lys Lys Arg Arg Gly
85 90 95

Arg Ala Thr Asp Ser Leu Pro Gly Lys Phe Glu Asp Met Tyr Lys Leu
100 105 110

Thr Ser Glu Leu Leu Gly Glu Gly Ala Tyr Ala Lys Val Gln Gly Ala
115 120 125

Val Ser Leu Gln Asn Gly Lys Glu Tyr Ala Val Lys Val Ser Val Ser

130

135

140

Ala Glu Cys Gln Ala Leu Leu Cys Lys
145 150

<210> 169

<211> 231

<212> PRT

<213> Homo sapiens

<400> 169

Met Gly Ser Gly Met Lys Leu Asn Asn Ser Cys Thr Pro Ile Thr Thr
1 5 10 15

Pro Glu Leu Thr Thr Pro Cys Gly Ser Ala Glu Tyr Met Ala Pro Glu
20 25 30

Val Val Glu Val Phe Thr Asp Gln Ala Thr Phe Tyr Asp Lys Arg Cys
35 40 45

Asp Leu Trp Ser Leu Gly Val Val Leu Tyr Ile Met Leu Ser Gly Tyr
50 55 60

Pro Pro Phe Val Gly His Cys Gly Ala Asp Cys Gly Trp Asp Arg Gly
65 70 75 80

Glu Val Cys Arg Val Cys Gln Asn Lys Leu Phe Glu Ser Ile Gln Glu
85 90 95

Gly Lys Tyr Glu Phe Pro Asp Lys Asp Trp Ala His Ile Ser Ser Glu
100 105 110

Ala Lys Asp Leu Ile Ser Lys Leu Leu Val Arg Asp Ala Lys Gln Arg
115 120 125

Leu Ser Ala Ala Gln Val Leu Gln His Pro Trp Val Gln Gly Gln Ala
130 135 140

Pro Glu Lys Gly Leu Pro Thr Pro Gln Val Leu Gln Arg Asn Ser Ser
145 150 155 160

Thr Met Asp Leu Thr Leu Phe Ala Ala Glu Ala Ile Ala Leu Asn Arg
165 170 175

Gln Leu Ser Gln His Glu Glu Asn Glu Leu Ala Glu Glu Pro Glu Ala
180 185 190

Leu Ala Asp Gly Leu Cys Ser Met Lys Leu Ser Pro Pro Cys Lys Ser
195 200 205

Arg Leu Ala Arg Arg Arg Ala Leu Ala Gln Ala Gly Arg Gly Glu Asp
210 215 220

Arg Ser Pro Pro Thr Ala Leu
225 230

<210> 170
<211> 146
<212> PRT
<213> Homo sapiens

<400> 170

Met Arg Lys Gly Val Leu Lys Asp Pro Glu Ile Ala Asp Leu Phe Tyr
1 5 10 15

Lys Asp Asp Pro Glu Glu Leu Phe Ile Gly Leu His Glu Ile Gly His
20 25 30

Gly Ser Phe Gly Ala Val Tyr Phe Ala Thr Asn Ala His Thr Ser Glu
35 40 45

Val Val Ala Ile Lys Lys Met Ser Tyr Ser Gly Lys Gln Thr His Glu
50 55 60

Lys Trp Gln Asp Ile Leu Lys Glu Val Lys Phe Leu Arg Gln Leu Lys
65 70 75 80

His Pro Asn Thr Ile Glu Tyr Lys Gly Cys Tyr Leu Lys Glu His Thr
85 90 95

Ala Trp Leu Val Met Glu Tyr Cys Leu Gly Ser Ala Ser Asp Leu Leu
100 105 110

Glu Val His Lys Lys Pro Leu Gln Glu Val Glu Ile Ala Ala Ile Thr
115 120 125

His Gly Ala Leu His Gly Leu Ala Tyr Leu His Ser His Ala Leu Ile
130 135 140

His Arg

145

<210> 171
<211> 123
<212> PRT
<213> Homo sapiens

<400> 171

Met Met Glu Glu Leu His Ser Leu Asp Pro Arg Arg Gln Glu Leu Leu
1 5 10 15

Glu Ala Arg Phe Thr Gly Val Gly Val Ser Lys Gly Pro Leu Asn Ser
20 25 30

Glu Ser Ser Asn Gln Ser Leu Cys Ser Val Gly Ser Leu Ser Asp Lys
35 40 45

Glu Val Glu Thr Pro Glu Lys Lys Gln Asn Asp Gln Arg Asn Arg Lys
50 55 60

Arg Lys Ala Glu Pro Tyr Glu Thr Ser Gln Gly Lys Gly Thr Pro Arg

65

70

75

80

Gly His Lys Ile Ser Asp Tyr Phe Glu Thr Ala Pro Leu Trp Phe Arg
85 90 95

Trp Gln Cys Cys Lys Gly Gly Asn Arg Gly Ala Val Cys Ser Ala Asn
100 105 110

Pro His Val Ser Asp Ala Ser Lys Thr Ser Ala
115 120

<210> 172

<211> 478

<212> PRT

<213> Homo sapiens

<400> 172

Met Val Gly Ile Lys Glu Arg Pro Ser Ser Asn Leu Pro Cys Pro Pro
1 5 10 15

Leu Pro Pro Gln Thr Gln Ala Cys Pro Pro Leu Ser Trp Pro Gln Arg
20 25 30

Leu Asp Ile Leu Leu Gly Thr Ala Arg Ala Ile Gln Phe Leu His Gln
35 40 45

Asp Ser Pro Ser Leu Ile His Gly Asp Ile Lys Ser Ser Asn Val Leu
50 55 60

Leu Asp Glu Arg Leu Thr Pro Lys Leu Gly Asp Phe Gly Leu Ala Arg
65 70 75 80

Phe Ser Arg Phe Ala Gly Ser Ser Pro Ser Gln Ser Ser Met Val Ala
85 90 95

Arg Thr Gln Thr Val Arg Gly Thr Leu Ala Tyr Leu Pro Glu Glu Tyr
100 105 110

Ile Lys Thr Gly Arg Leu Ala Val Asp Thr Asp Thr Phe Ser Phe Gly
115 120 125

Val Val Val Leu Glu Thr Leu Ala Gly Gln Arg Ala Val Lys Thr His
130 135 140

Gly Ala Arg Thr Lys Tyr Leu Lys Asp Leu Val Glu Glu Ala Glu
145 150 155 160

Glu Ala Gly Val Ala Leu Arg Ser Thr Gln Ser Thr Leu Gln Ala Gly
165 170 175

Leu Ala Ala Asp Ala Trp Ala Ala Pro Ile Ala Met Gln Ile Tyr Lys
180 185 190

Lys His Leu Asp Pro Arg Pro Gly Pro Cys His Leu Ser Trp Ala Trp
195 200 205

Ala Trp Ala Ser Trp Pro Ala Ala Ala Cys Thr Ala Gly Pro Lys Gly
210 215 220

Arg Pro Pro Met Thr Gln Val Tyr Glu Arg Leu Glu Lys Leu Gln Ala
225 230 235 240

Val Val Ala Gly Val Pro Gly His Leu Glu Ala Ala Ser Cys Ile Pro
245 250 255

Phe Pro Gln Glu Asn Ser Tyr Val Ser Ser Thr Gly Arg Ala His Ser
260 265 270

Gly Ala Ala Pro Trp Gln Pro Leu Ala Ala Pro Ser Gly Ala Ser Ala
275 280 285

Gln Ala Ala Glu Gln Leu Gln Arg Gly Pro Asn Gln Pro Val Glu Ser
290 295 300

Asp Glu Ser Leu Gly Gly Leu Ser Ala Ala Leu Arg Ser Trp His Leu
305 310 315 320

Thr Pro Ser Cys Pro Leu Asp Pro Ala Pro Leu Arg Glu Ala Gly Cys
325 330 335

Pro Gln Gly Asp Thr Ala Gly Glu Ser Ser Trp Gly Ser Gly Pro Gly
340 345 350

Ser Arg Pro Thr Ala Val Glu Gly Leu Ala Leu Gly Ser Ser Ala Ser
355 360 365

Ser Ser Ser Glu Pro Pro Gln Ile Ile Ile Asn Pro Ala Arg Gln Lys
370 375 380

Met Val Gln Lys Leu Ala Leu Tyr Glu Asp Gly Ala Leu Asp Ser Leu
385 390 395 400

Gln Leu Leu Ser Ser Ser Leu Pro Gly Leu Gly Leu Glu Gln Asp
405 410 415

Arg Gln Gly Pro Lys Lys Val Met Asn Phe Arg Ala Asp Val Phe Thr
420 425 430

Trp Ala Asp Pro Pro Asn Pro Glu Val Lys Val Leu Met Val Arg Ser
435 440 445

Ser His Gly Ala Arg Val Leu Ser Thr Leu Pro Ala Val Gly Val Gly
450 455 460

Ala His Ala Arg Trp Gly Glu Lys Glu Val Ala Leu Leu Phe
465 470 475

<210> 173
<211> 344
<212> PRT
<213> Homo sapiens

<400> 173

Met Ala Gly Gly Pro Gly Pro Gly Glu Pro Ala Ala Pro Gly Ala Gln
1 5 10 15

His Phe Leu Tyr Glu Val Pro Pro Trp Val Met Cys Arg Phe Tyr Lys
20 25 30

Val Met Asp Ala Leu Glu Pro Ala Asp Trp Cys Gln Phe Ala Ala Leu
35 40 45

Ile Val Arg Asp Gln Thr Glu Leu Arg Leu Cys Glu Arg Ser Gly Gln
50 55 60

Arg Thr Ala Ser Val Leu Trp Pro Trp Ile Asn Arg Asn Ala Arg Val
65 70 75 80

Ala Asp Leu Val His Ile Leu Thr His Leu Gln Leu Leu Arg Ala Arg
85 90 95

Asp Ile Ile Thr Ala Trp His Pro Pro Ala Pro Leu Pro Ser Pro Gly
100 105 110

Thr Thr Ala Pro Arg Pro Ser Ser Ile Pro Ala Pro Ala Glu Ala Glu
115 120 125

Ala Trp Ser Pro Arg Lys Leu Pro Ser Ser Ala Ser Thr Phe Leu Ser
130 135 140

Pro Ala Phe Pro Gly Ser Gln Thr His Ser Gly Pro Glu Leu Gly Leu
145 150 155 160

Val Pro Ser Pro Ala Ser Leu Trp Pro Pro Pro Ser Pro Ala Pro
165 170 175

Ser Ser Thr Lys Pro Gly Pro Glu Ser Ser Val Ser Leu Leu Gln Gly
180 185 190

Ala Arg Pro Ser Pro Phe Cys Trp Pro Leu Cys Glu Ile Ser Arg Gly
195 200 205

Thr His Asn Phe Ser Glu Glu Leu Lys Ile Gly Glu Gly Phe Gly
210 215 220

Cys Val Tyr Arg Ala Val Met Arg Asn Thr Val Tyr Ala Val Lys Arg
225 230 235 240

Leu Lys Glu Asn Ala Asp Leu Glu Trp Thr Ala Val Lys Gln Ser Phe
245 250 255

Leu Thr Glu Val Glu Gln Leu Ser Arg Phe Arg His Pro Asn Ile Val
260 265 270

Asp Phe Ala Gly Tyr Cys Ala Gln Asn Gly Phe Tyr Cys Leu Val Tyr
275 280 285

Gly Phe Leu Pro Asn Gly Ser Leu Glu Asp Arg Leu His Cys Gln Thr
290 295 300

Gln Ala Cys Pro Pro Leu Ser Trp Pro Gln Arg Leu Asp Ile Leu Leu
305 310 315 320

Gly Thr Ala Arg Ala Ser Gln Val Ser Cys Asn Arg Val Ser Ser Cys
325 330 335

Val Ser Lys Ser Ser Pro Gly Leu
340

<210> 174
<211> 336
<212> PRT
<213> Homo sapiens

<400> 174

Met Phe Thr Glu Glu Asp Val Lys Phe Tyr Leu Ala Glu Leu Ala Leu
1 5 10 15

Ala Leu Asp His Leu His Ser Leu Gly Ile Ile Tyr Arg Asp Leu Lys
20 25 30

Pro Glu Asn Ile Leu Leu Asp Glu Glu Gly His Ile Lys Leu Thr Asp
35 40 45

Phe Gly Leu Ser Lys Glu Ser Ile Asp His Glu Lys Lys Ala Tyr Ser
50 55 60

Phe Cys Gly Thr Val Glu Tyr Met Ala Pro Glu Val Val Asn Arg Arg
65 70 75 80

Gly His Thr Gln Ser Ala Asp Trp Trp Ser Phe Gly Val Leu Met Phe
85 90 95

Glu Met Leu Thr Gly Thr Leu Pro Phe Gln Gly Lys Asp Arg Lys Glu
100 105 110

Thr Met Thr Met Ile Leu Lys Ala Lys Leu Gly Met Pro Gln Phe Leu
115 120 125

Ser Pro Glu Ala Gln Ser Leu Leu Arg Met Leu Phe Lys Arg Asn Pro
130 135 140

Ala Asn Arg Leu Gly Ala Gly Pro Asp Gly Val Glu Glu Ile Lys Arg
145 150 155 160

His Ser Phe Phe Ser Thr Ile Asp Trp Asn Lys Leu Tyr Arg Arg Glu
165 170 175

Ile His Pro Pro Phe Lys Pro Ala Thr Gly Arg Pro Glu Asp Thr Phe
180 185 190

Tyr Phe Asp Pro Glu Phe Thr Ala Lys Thr Pro Lys Asp Ser Pro Gly
195 200 205

Ile Pro Pro Ser Ala Asn Ala His Gln Leu Phe Arg Gly Phe Ser Phe

210

215

220

Val Ala Ile Thr Ser Asp Asp Glu Ser Gln Ala Met Gln Thr Val Gly
225 230 235 240

Val His Ser Ile Val Gln Gln Leu His Arg Asn Ser Ile Gln Phe Thr
245 250 255

Asp Gly Tyr Glu Val Lys Glu Asp Ile Gly Val Gly Ser Tyr Ser Val
260 265 270

Cys Lys Arg Cys Ile His Lys Ala Thr Asn Met Glu Phe Ala Val Lys
275 280 285

Val Asn Phe Phe Tyr Leu Lys Cys Asn Ser Tyr Ser Ser Cys Ser Cys
290 295 300

Met Ser Val Pro Val Lys Asn Tyr Thr Pro Leu Val Val Lys Ser Ala
305 310 315 320

Phe Cys Tyr Lys Lys Val Lys Tyr Leu Ala Ser Asp Leu Gln Arg Ser
325 330 335

<210> 175

<211> 198

<212> PRT

<213> Homo sapiens

<400> 175

Met Pro Leu Ala Gln Leu Ala Asp Pro Trp Gln Lys Met Ala Val Glu
1 5 10 15

Ser Pro Ser Asp Ser Ala Glu Asn Gly Gln Gln Ile Met Asp Glu Pro
20 25 30

Met Gly Glu Glu Glu Ile Asn Pro Gln Thr Glu Glu Val Ser Ile Lys
35 40 45

Glu Ile Ala Ile Thr His His Val Lys Glu Gly His Glu Lys Ala Asp
50 55 60

Pro Ser Gln Phe Glu Leu Leu Lys Val Leu Gly Gln Gly Ser Phe Gly
65 70 75 80

Lys Val Phe Leu Val Lys Lys Ile Ser Gly Ser Asp Ala Arg Gln Leu
85 90 95

Tyr Ala Met Lys Val Leu Lys Lys Ala Thr Leu Lys Val Arg Asp Arg
100 105 110

Val Arg Thr Lys Met Glu Arg Asp Ile Leu Val Glu Val Asn His Pro
115 120 125

Phe Ile Val Lys Leu His Tyr Ala Phe Gln Thr Glu Gly Lys Leu Tyr
130 135 140

Leu Ile Leu Asp Phe Leu Arg Gly Gly Asp Leu Phe Thr Arg Leu Ser
145 150 155 160

Lys Glu Val Met Phe Thr Glu Glu Asp Val Lys Phe Tyr Leu Ala Glu
165 170 175

Leu Ala Leu Ala Leu Asp His Leu His Ser Leu Gly Ile Ile Tyr Arg
180 185 190

Asp Leu Lys Pro Glu Lys
195

<210> 176

<211> 489

<212> PRT

<213> Homo sapiens

<400> 176

Met Ser Thr Glu Ala Asp Glu Gly Ile Thr Phe Ser Val Pro Pro Phe
1 5 10 15

Ala Pro Ser Gly Phe Cys Thr Ile Pro Glu Gly Gly Ile Cys Arg Arg
20 25 30

Gly Gly Ala Ala Ala Val Gly Glu Gly Glu His Gln Leu Pro Pro
35 40 45

Pro Pro Pro Gly Ser Phe Trp Asn Val Glu Ser Ala Ala Ala Pro Gly
50 55 60

Ile Gly Cys Pro Ala Ala Thr Ser Ser Ser Ala Thr Arg Gly Arg
65 70 75 80

Gly Ser Ser Val Gly Gly Ser Arg Arg Thr Thr Val Ala Tyr Val
85 90 95

Ile Asn Glu Ala Ser Gln Gly Gln Leu Val Val Ala Glu Ser Glu Ala
100 105 110

Leu Gln Ser Leu Arg Glu Ala Cys Glu Thr Val Gly Ala Thr Leu Glu
115 120 125

Thr Leu His Phe Gly Lys Leu Asp Phe Gly Glu Thr Thr Val Leu Asp
130 135 140

Arg Phe Tyr Asn Ala Asp Ile Ala Val Val Glu Met Ser Asp Ala Phe
145 150 155 160

Arg Gln Pro Ser Leu Phe Tyr His Leu Gly Val Arg Glu Ser Phe Ser
165 170 175

Met Ala Asn Asn Ile Ile Leu Tyr Cys Asp Thr Asn Ser Asp Ser Leu
180 185 190

Gln Ser Leu Lys Glu Ile Ile Cys Gln Lys Asn Thr Met Cys Thr Gly
195 200 205

Asn Tyr Thr Phe Val Pro Tyr Met Ile Thr Pro His Asn Lys Val Tyr
210 215 220

Cys Cys Asp Ser Ser Phe Met Lys Gly Leu Thr Glu Leu Met Gln Pro
225 230 235 240

Asn Phe Glu Leu Leu Leu Gly Pro Ile Cys Leu Pro Leu Val Asp Arg
245 250 255

Phe Ile Gln Leu Leu Lys Val Ala Gln Ala Ser Ser Ser Gln Tyr Phe
260 265 270

Arg Glu Ser Ile Leu Asn Asp Ile Arg Lys Ala Arg Asn Leu Tyr Thr
275 280 285

Gly Lys Glu Leu Ala Ala Glu Leu Ala Arg Ile Arg Gln Arg Val Asp
290 295 300

Asn Ile Glu Val Leu Thr Ala Asp Ile Val Ile Asn Leu Leu Leu Ser
305 310 315 320

Tyr Arg Asp Ile Gln Asp Tyr Asp Ser Ile Val Lys Leu Val Glu Thr
325 330 335

Leu Glu Lys Leu Pro Thr Phe Asp Leu Ala Ser His His Val Lys
340 345 350

Phe His Tyr Ala Phe Ala Leu Asn Arg Arg Asn Leu Pro Gly Asp Arg
355 360 365

Ala Lys Ala Leu Asp Ile Met Ile Pro Met Val Gln Ser Glu Gly Gln
370 375 380

Val Ala Ser Asp Met Tyr Cys Leu Val Gly Arg Ile Tyr Lys Asp Met
385 390 395 400

Phe Leu Asp Ser Asn Phe Thr Asp Thr Glu Ser Arg Asp His Gly Ala
405 410 415

Ser Trp Phe Lys Lys Ala Phe Glu Ser Glu Pro Thr Leu Gln Ser Gly
420 425 430

Ile Asn Tyr Ala Val Leu Leu Leu Ala Ala Gly His Gln Phe Glu Ser
435 440 445

Ser Phe Glu Leu Arg Lys Val Gly Asn Tyr Asn Leu Asn Phe Tyr Met
450 455 460

Glu Ile Lys Lys Leu Gly Pro Asn Leu Val Gln Arg Arg Ile Ser Ala
465 470 475 480

Asp Ser Asp Gly Ser Pro Gly Phe Val
485

<210> 177

<211> 105

<212> PRT
<213> Homo sapiens

<400> 177

Met Arg Glu Phe Glu Val Leu Lys Lys Leu Asn His Lys Asn Ile Val
1 5 10 15

Lys Leu Phe Ala Ile Glu Glu Glu Thr Thr Arg His Lys Val Leu
20 25 30

Ile Met Glu Phe Cys Pro Cys Gly Ser Leu Tyr Thr Val Leu Glu Glu
35 40 45

Pro Ser Asn Ala Tyr Gly Leu Pro Glu Ser Glu Phe Leu Ile Val Leu
50 55 60

Arg Asp Val Val Gly Gly Met Asn His Leu Arg Glu Asn Gly Ile Val
65 70 75 80

His Arg Asp Ile Lys Pro Gly Asn Ile Met Arg Ala Leu Tyr His Ser
85 90 95

Leu Val Asp Asp Ser Phe His Pro Pro
100 105

<210> 178

<211> 413

<212> PRT

<213> Homo sapiens

<220>

<221> -

<222> (1)..(413)

<223> "XAA" can be any amino acid

<400> 178

Met Tyr Cys Phe Gly Arg Lys Xaa Tyr Ile Ser Thr Arg Pro Cys Phe
1 5 10 15

Pro Asn Lys Thr Cys Gln Lys Met Leu Ile Ile Leu Thr Ser Ala Leu
20 25 30

Gln Ile Ala His Arg Cys Ile Cys Arg Ile Leu Leu Gly Ser Arg Val
35 40 45

Leu Ala Ala Lys Ala Ser Gly Asn Cys Thr Leu Asn Ser Glu Asp Phe
50 55 60

Ile Phe Asn Ile Gly Ser Ala Ala Tyr Asp Ala Val Leu Asp Arg Asn
65 70 75 80

Val Ala Ile Lys Lys Leu Ser Arg Pro Phe Gln Asn Gln Thr His Ala
85 90 95

Lys Arg Ala Tyr Arg Glu Leu Val Leu Met Lys Cys Val Asn His Lys
100 105 110

Asn Ile Ile Ser Leu Leu Asn Val Phe Thr Pro Gln Lys Thr Leu Glu
115 120 125

Glu Phe Gln Asp Val Tyr Leu Val Met Glu Leu Met Asp Ala Asn Leu
130 135 140

Cys Gln Val Ile Gln Met Glu Leu Asp His Glu Arg Met Ser Tyr Leu
145 150 155 160

Leu Tyr Gln Met Leu Cys Gly Ile Lys His Leu His Ser Ala Gly Ile
165 170 175

Ile His Arg Asp Leu Lys Pro Ser Asn Ile Val Val Lys Ser Asp Cys
180 185 190

Thr Leu Lys Ile Leu Asp Phe Gly Leu Ala Arg Thr Ala Gly Thr Ser
195 200 205

Phe Met Met Thr Pro Tyr Val Val Thr Arg Tyr Tyr Arg Ala Pro Glu
210 215 220

Val Ile Leu Gly Met Gly Tyr Lys Glu Asn Val Asp Ile Trp Ser Val
225 230 235 240

Gly Cys Ile Met Gly Glu Met Val Arg His Lys Ile Leu Phe Pro Gly
245 250 255

Arg Asp Tyr Ile Asp Gln Trp Asn Lys Val Ile Glu Gln Leu Gly Thr
260 265 270

Pro Cys Pro Glu Phe Met Lys Lys Leu Gln Pro Thr Val Arg Asn Tyr
275 280 285

Val Glu Asn Arg Pro Lys Tyr Ala Gly Leu Thr Phe Pro Lys Leu Phe
290 295 300

Pro Asp Ser Leu Phe Pro Ala Asp Ser Glu His Asn Lys Leu Lys Ala
305 310 315 320

Ser Gln Ala Arg Asp Leu Leu Ser Lys Met Leu Val Ile Asp Pro Ala
325 330 335

Lys Arg Ile Ser Val Asp Asp Ala Leu Gln His Pro Tyr Ile Asn Val
340 345 350

Trp Tyr Asp Pro Ala Glu Val Glu Ala Pro Pro Pro Gln Ile Tyr Asp
355 360 365

Lys Gln Leu Asp Glu Arg Glu His Thr Ile Glu Glu Trp Lys Glu Leu
370 375 380

Ile Tyr Lys Glu Val Met Asn Ser Glu Glu Lys Thr Lys Asn Gly Val
385 390 395 400

Val Lys Gly Gln Pro Ser Pro Ser Ala Gln Val Gln Gln
405 410

<210> 179
<211> 108
<212> PRT
<213> Homo sapiens

<400> 179

Met Ser Lys Ser Lys Val Asp Asn Gln Phe Tyr Ser Val Glu Val Gly
1 5 10 15

Asp Ser Thr Phe Thr Val Leu Lys Arg Tyr Gln Asn Leu Lys Pro Ile
20 25 30

Gly Ser Gly Ala Gln Gly Ile Val Cys Ala Ala Tyr Asp Ala Val Leu
35 40 45

Asp Arg Asn Val Ala Ile Lys Lys Leu Ser Arg Pro Phe Gln Asn Gln
50 55 60

Thr His Ala Lys Arg Ala Tyr Arg Glu Leu Val Leu Met Lys Cys Val
65 70 75 80

Asn His Lys Asn Val Ser Phe Val Ile Phe Lys Leu Leu Ala Val Gly
85 90 95

Val Cys Lys Ile Gly Lys Arg Lys Cys Val Cys Thr
100 105

<210> 180
<211> 336
<212> PRT
<213> Homo sapiens

<400> 180

Met Ala Met Thr Gly Ser Thr Pro Cys Ser Ser Met Ser Asn His Thr
1 5 10 15

Lys Glu Arg Val Thr Met Thr Lys Val Thr Leu Glu Asn Phe Tyr Ser
20 25 30

Asn Leu Ile Ala Gln His Glu Glu Arg Glu Met Arg Gln Lys Lys Leu
35 40 45

Glu Lys Val Met Glu Glu Glu Gly Leu Lys Asp Glu Glu Lys Arg Leu
50 55 60

Arg Arg Ser Ala His Ala Arg Lys Glu Thr Glu Phe Leu Arg Leu Lys
65 70 75 80

Arg Thr Arg Leu Gly Leu Glu Asp Phe Glu Ser Leu Lys Val Ile Gly
85 90 95

Arg Gly Ala Phe Gly Glu Val Arg Leu Val Gln Lys Lys Asp Thr Gly

100

105

110

His Val Tyr Ala Met Lys Ile Leu Arg Lys Ala Asp Met Leu Glu Lys
115 120 125

Glu Gln Val Gly His Ile Arg Ala Glu Arg Asp Ile Leu Val Glu Ala
130 135 140

Asp Ser Leu Trp Val Val Lys Met Phe Tyr Ser Phe Gln Asp Lys Leu
145 150 155 160

Asn Leu Tyr Leu Ile Met Glu Phe Leu Pro Gly Gly Asp Met Met Thr
165 170 175

Leu Leu Met Lys Lys Asp Thr Leu Thr Glu Glu Glu Thr Gln Phe Tyr
180 185 190

Ile Ala Glu Thr Val Leu Ala Ile Asp Ser Ile His Gln Leu Gly Phe
195 200 205

Ile His Arg Asp Ile Lys Pro Asp Asn Leu Leu Asp Ser Lys Gly
210 215 220

His Val Lys Leu Ser Asp Phe Gly Leu Cys Thr Gly Leu Lys Lys Ala
225 230 235 240

His Arg Thr Glu Phe Tyr Arg Asn Leu Asn His Ser Leu Pro Ser Asp
245 250 255

Phe Thr Phe Gln Asn Met Asn Ser Lys Arg Lys Ala Glu Thr Trp Lys
260 265 270

Arg Asn Arg Arg Gln Leu Ala Phe Ser Thr Val Gly Thr Pro Asp Tyr
275 280 285

Ile Ala Pro Glu Val Phe Met Gln Thr Gly Tyr Asn Lys Leu Cys Asp
290 295 300

Trp Trp Ser Leu Gly Val Ile Met Tyr Glu Met Leu Ile Gly Lys Leu
305 310 315 320

His Gly Phe Arg Gly Leu Phe Leu Cys Ile His Asp Arg Leu Leu His
325 330 335

<210> 181

<211> 415

<212> PRT

<213> Homo sapiens

<220>

<221> -

<222> (1)..(415)

<223> "XAA " can be any amino acid

<400> 181

Xaa Arg His Glu Ser Ala Arg Ala Ala Arg Val Ser Gly Gly Ser Met
1 5 10 15

Leu Asp Ile Ile Lys Tyr Ile Val Asn Arg Gly Glu His Lys Asn Gly
20 25 30

Val Leu Glu Glu Ala Ile Ile Ala Thr Ile Leu Lys Glu Val Leu Glu
35 40 45

Gly Leu Asp Tyr Leu His Arg Asn Gly Gln Ile His Arg Asp Leu Lys
50 55 60

Ala Gly Asn Ile Leu Leu Gly Glu Asp Gly Ser Val Gln Ile Ala Asp
65 70 75 80

Phe Gly Val Ser Ala Phe Leu Ala Thr Gly Gly Asp Val Thr Arg Asn
85 90 95

Lys Val Arg Lys Thr Phe Val Gly Thr Pro Cys Trp Met Ala Pro Glu
100 105 110

Val Met Glu Gln Val Arg Gly Tyr Asp Phe Lys Ala Asp Met Trp Ser
115 120 125

Phe Gly Ile Thr Ala Ile Glu Leu Ala Thr Gly Ala Ala Pro Tyr His
130 135 140

Lys Tyr Pro Pro Met Lys Val Leu Met Leu Thr Leu Gln Asn Asp Pro
145 150 155 160

Pro Thr Leu Glu Thr Gly Val Glu Asp Lys Glu Met Met Lys Lys Tyr
165 170 175

Gly Lys Ser Phe Arg Lys Leu Leu Ser Leu Cys Leu Gln Lys Asp Pro
180 185 190

Ser Lys Arg Pro Thr Ala Ala Glu Leu Leu Lys Cys Lys Phe Phe Gln
195 200 205

Lys Ala Lys Asn Arg Glu Tyr Leu Ile Glu Lys Leu Leu Thr Arg Thr
210 215 220

Pro Asp Ile Ala Gln Arg Ala Lys Lys Val Arg Arg Val Pro Gly Ser
225 230 235 240

Ser Gly His Leu His Lys Thr Glu Asp Gly Asp Trp Glu Trp Ser Asp
245 250 255

Asp Glu Met Asp Glu Lys Ser Glu Glu Gly Lys Ala Ala Phe Ser Gln
260 265 270

Glu Lys Ser Arg Arg Val Lys Glu Glu Asn Pro Glu Ile Ala Val Ser
275 280 285

Ala Ser Thr Ile Pro Glu Gln Ile Gln Ser Leu Ser Val His Asp Ser
290 295 300

Gln Gly Pro Pro Asn Ala Asn Glu Asp Tyr Arg Glu Ala Ser Ser Cys
305 310 315 320

Ala Val Asn Leu Val Leu Arg Leu Arg Asn Ser Arg Lys Glu Leu Asn
325 330 335

Asp Ile Arg Phe Glu Phe Thr Pro Gly Arg Asp Thr Ala Asp Gly Val
340 345 350

Ser Gln Glu Leu Phe Ser Ala Gly Leu Val Asp Gly His Asp Val Val
355 360 365

Ile Val Ala Ala Asn Leu Gln Lys Ile Val Asp Asp Pro Lys Ala Leu
370 375 380

Lys Thr Leu Thr Phe Lys Leu Ala Ser Gly Cys Asp Gly Ser Glu Ile
385 390 395 400

Pro Asp Glu Val Lys Leu Ile Gly Phe Ala Gln Leu Ser Val Ser
405 410 415

<210> 182

<211> 409

<212> PRT

<213> Homo sapiens

<220>

<221> -

<222> (1)..(409)

<223> "Xaa" can be any amino acid

<400> 182

Xaa Arg His Glu Ser Ala Arg Ala Ala Arg Val Ser Gly Gly Ser Met
1 5 10 15

Leu Asp Ile Ile Lys Tyr Ile Val Asn Arg Gly Glu His Lys Asn Gly
20 25 30

Val Leu Glu Glu Ala Ile Ile Ala Thr Ile Leu Lys Glu Val Leu Glu
35 40 45

Gly Leu Asp Tyr Leu His Arg Asn Gly Gln Ile His Arg Asp Leu Lys
50 55 60

Ala Gly Asn Ile Leu Leu Gly Glu Asp Gly Ser Val Gln Ile Ala Asp
65 70 75 80

Phe Gly Val Ser Ala Phe Leu Ala Thr Gly Gly Asp Val Thr Arg Asn
85 90 95

Lys Val Arg Lys Thr Phe Val Gly Thr Pro Cys Trp Met Ala Pro Glu
100 105 110

Val Met Glu Gln Val Arg Gly Tyr Asp Phe Lys Ala Asp Met Trp Ser
115 120 125

Phe Gly Ile Thr Ala Ile Glu Leu Ala Thr Gly Ala Ala Pro Tyr His
130 135 140

Lys Tyr Pro Pro Met Lys Val Leu Met Leu Thr Leu Gln Asn Asp Pro
145 150 155 160

Pro Thr Leu Glu Thr Gly Val Glu Asp Lys Glu Met Met Lys Lys Tyr
165 170 175

Gly Lys Ser Phe Arg Lys Leu Leu Ser Leu Cys Leu Gln Lys Asp Pro
180 185 190

Ser Lys Arg Pro Thr Ala Ala Glu Leu Leu Lys Cys Lys Phe Phe Gln
195 200 205

Lys Ala Lys Asn Arg Glu Tyr Leu Ile Glu Lys Leu Leu Thr Arg Thr
210 215 220

Pro Asp Ile Ala Gln Arg Ala Lys Lys Val Arg Arg Val Pro Gly Ser
225 230 235 240

Ser Gly His Leu His Lys Thr Glu Asp Gly Asp Trp Glu Trp Ser Asp
245 250 255

Asp Glu Met Asp Glu Lys Ser Glu Glu Gly Lys Ala Ala Phe Ser Gln
260 265 270

Glu Lys Ser Arg Arg Val Lys Glu Glu Asn Pro Glu Ile Ala Val Ser
275 280 285

Ala Ser Thr Ile Pro Glu Gln Ile Gln Ser Leu Ser Val His Asp Ser
290 295 300

Gln Gly Pro Pro Asn Ala Asn Glu Asp Tyr Arg Glu Ala Ser Ser Cys
305 310 315 320

Ala Val Asn Leu Val Leu Arg Leu Arg Asn Ser Arg Lys Glu Leu Asn
325 330 335

Asp Ile Arg Phe Glu Phe Thr Pro Gly Arg Asp Thr Ala Asp Gly Val
340 345 350

Ser Gln Glu Leu Phe Ser Ala Gly Leu Val Asp Gly His Asp Val Val
355 360 365

Ile Val Ala Ala Asn Leu Gln Lys Ile Val Asp Asp Pro Lys Ala Leu
370 375 380

Lys Thr Leu Thr Phe Lys Leu Asn Gln Phe Leu His Leu Glu Ala Phe
385 390 395 400

Asp Ser Ala Ala Leu Gly Asn Val Phe
405